>SGPR397 SEQID

FIG. 1A

>SGPR413 SEQID 2

ATTGGTCCTGCCTTTTGTCAGTGGTTTGTAAAAGAAGCTCTTCTAACATATAAGAGTGACCCAGCCATGAGAAAA TGCTGAATCATCTATATTTCTATATCATGCCTGTGTTTAACGTCGATGGATACCATTTTAGTTGGACCAATGATCGA AGAGGAAGCATATGCACTGAAGAAAATATCCTATCAACTTAAGGTGGACCTGTGGCAGCCCAGCAGTATCTCCTA ATITTAAAGCTGGGCAGACGATCACGACTCAAAAGAGCTGTTTGGATAGACTGTGGGTATTCATGCAAGAGAATGG GCAACCGGGGCACAGCCACCTTTATAACAACCGCTATGCTGGTGATAAAGTGATAAGATTTATTCCCAAAACAGA TGTATCAGAGGGAACAGTTACTGATGTCCATATCCCCCAAAATGGTTCCCGAGCCCTGTTAGCCTTCTTACAGGA AGCCAACATCCAGTACAAGGTCCTCATAGAAGATCTTCAGAAAACACTGGAGAAGGGAAGCAGCTTGCACACCC AGAGAAACCGAAGATCCCTCTGGATATAATTATGAAGTTTATCACTCCTTAGAAGAAATTCAAAATTGGATGCA GAAGTGGTGGTAAGTTTGGGACCAACTGGGATCCAGATCCAAAGGTTTCTGCAGGTTTTACTCTGCAAAATAT TTTGGAGAAAAACAAGGTCAAGGAACTCAAGGTTTCGCTGCCGTGGAGTGGATGCCAATAGAAACTGGAAAGT ATGAAGTGTCTCGGGAAGCGCAGGGGCCAGGCAGCTGCTTTCCTGCCTCTTTGCTGGCTCTTTTGAAGATTCT GAGTCCAGAGGACTCTCATGGGAGACTCATGTTTTCTGTATGTGA

CATGTTTTGGGACGGATCTCAATCGAAATTTCAATGCATCTTGGTGTAGTATTGGTGCCTCTAGAAACTGCCAAGA GCCCAACACAGAGAGAGATTGTGGACAAGTCAGTGAGTCCATGGAGCCTGGAGACGTATTCCTATAACATATAC AACATAGATGGTTATATCTACACTTGGACAACTGATCGTCTTTGGAGGAAATCCCGTTCACCCCATAATAATGGCA 4GGATGATATTTTGTGCTTCCTGACCATGCACTCTTATGGGCAGTTAATTCTCACACCTTACGGCTACACAAAAA SCAGCCCACCTGTGAGGAGCCATGGAGGCTGTGCTGTCAGTCCTGGATGATGTGTATGCGAAACACTGGCACT -AAATCAAGTAACCACCCAGAAATGATTCAAGTTGGACAGAAGGCAGCAAATGCATTGAAAGCAAAGTATGGAAC 'GGGATTCCCTTCTCATATACGTTTGAGCTGAGGGACAGTGGAACATATGGGTTTGTTCTGCCAGAAGCTCAGAT ATGAAGCCTCTGCTTGAAACCCTTTATCTTTTGGGGATGCTGGTTCCTGGAGGGCTGGGATATGATAGATCCTTA SGGACAGTGCTGGAAGGGTGACATCTGCCACTATGCTGCTGGGCCTGCTGGTGTCCTGCATGTCTCTTCTTAA CCTAGGAGTGACCTATGAGACCCACCCCATGTATTATCTGAAGATCAGCCAACCATCTGGTAATCCCAAGAAAT ICAAACATTCTGTGGGACAGGGCCAGTGTCTGAACCAGAGACTAAAGCTGTTGCCAGCTTCATAGAGAGCAAGA CACCCCATGGGAGAGATCTATGAGTGGATGAGAGAGATCAGTGAGAAGTACAAGGAAGTGGTGACACAGCATTT CATTTGGATGGACTGTGGAATTCACGCCAGAGAATGGATTGCTCCTGCTTTTTGCCAATGGTTCGTCAAAGAAAT CAATTATAGAGTTGGATCGAGTGCAGATATTTTATATGCCTCATCAGGGTCTTCAAGAGATTGGGCCCGAGACAT ICTACAAAACCATAAAGACAACTCAAGTATACGCAAGCTCCTTAGGAACCTGGACTTCTATGTCCTTCCAGTTCT

>SGPR404 SEQID 3

FIG. 1B

GTTTCACTGCATCCACCAAGAACTGTATGGTTGGCTATGACATGGGGGCCACAAGGTGTGACTTCACACTTAGCA AGCTGTATGCTGTGGAGATCTCAGATCACCCTGGGGAGCATGAAGTCGGTGAGCCCGAGTTCCACTACATCGCG GAATTGACATCAACAACATTTCCTGATTTAAACACGCTGCTCTGGGAGGCAGAGGATCGACAGAATGTCCCCA ACGGAGGACTTCCAGAAGGAGGAGGACTGTCAATGGGGCCTCCTGGCACACCGTCGCTGGAAGTCTGAACG AAACCAACATGGCCAGGATCCGAGAGATCATGGAGAAGTTTGGGAAGCAGCCCGTCAGCCTGCCAGGCGG ATCACCGCCGGAACGAGATGACCACCACTGATGACCTGGATTTTAAGCACCACAATTATAAGGAAATGCGCCAGI ATGECTACGAGAGGCCTACGAAGGGGGCTCGGAGCTGGGAGGCTGGTCCCTGGGACGCTGGACCCACGATG AGAGCAGTCATAGCCTGGATGGAAAAAATCCCTTTTGTGCTGGGCGGCGACCTGCAGGGCGGCGGCGGTGGTGG TGGCGTACCCCTACGACCTGGTGCGGTCCCCCTGGAAGACGCAGGAACACACCCCCCACCCCCGACGACACG AGCTGCCCGAGGAGTGGGAGAATAACCGGGAATCTCTGATCGTGTTCATGGAGCAGGTTCATGGTGGCATTAAA GGCTTGGTGAGAGATTCACATGGAAAAGGAATCCCAAACGCCATTATCTCCGTAGAAGGCATTAACCATGACATC GTTCCGCTGGCTGGCCTACTCCTATGCCTCCACACACGCCTCATGACAGACGCCCGGAGGAGGGGGGTGTGCCAC CGAACAGCCAACGATGGGGGATTACTGGCGCCTCCTGAACCCTGGAGAGTATGTGGTCACAGCAAAGGCCGAAG ATGGTGAGCAATGACAGCCACACGTGGGTCACTGTTAAGAATGGATCTGGAGACATGATATTTGAGGGAAACAG CCTGGTTTGATAATGGGAGCATCTGCATGAGAATGGAGATCCTGGGCTGCCCACTGCCAGATCCTAATAATTATT TGATGAAAGTTGTGAATGAAATGTGTCCCAATATCACCAGAATTTACAACATTGGAAAAAGCCACCAGGGCCTGA GGGCCCACGGCAATGAGGTGCTGGGCCGGGAGCTGCTGCTGCTGGTGGTGCAGTTCGTGTCAGGAGTAC GGAAAGTTCCCAATCACTATATTGCAATCCCTGAGTGGTTTCTGTCGGAAAATGCCACGGTGGCTGCCGAGACC ATTTCAGCTACCTTCATACAAACTGCTTCGAACTGTCCATCTACGTGGGCTGTGATAAATACCCACATGAGAGCC TGAGAAGGAGATCCCTGTTCTCAATGAGCTACCCGTCCCCATGGTGGCCCGCTACATCCGCATAAACCCTCAGT GCTGAAGCTGCGGGGGGGGGAAGAGACGACAGCGTGGGTGA

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GTGACCTCTTCTGCAACCGCACGGTCTCCGACTGCTGCCTGACTTCTGGGACTTCTGCCTCGGCGTGCCACCC CAGGAGCAGGACCTGTGCTGCCGCGGCCGTGCCGACGACTGTGCCCTGCCCTACCTGGGCGCGCATCTGTTACT GTCGTGGGCGCCCGGGAGCTAGCACCGGGTCTGCACCTGCGGGGCATCCGGGACGCGGGAGGCCGGTACTGC AACCGTTGCACCTGCCAGGAGAACAGGCAGTGGCAGTGTGACCAAGAACCATGCCTGGTGGATCCAGACATGA TCAAAGCCATCAACCAGGGCAACTATGGCTGGCAGGCTGGGAACCACAGCGCCTTCTGGGGCATGACCCTGGA

CCTTCTCGGGCCGTGAACGAGACGAGGCTGGCCCTGCGCCCCCTGTATGATGCACAGCCGAGCCATGGGTCG TCTACCECCTCEGCTCCAACGACAAGGAGATCATGAAGGAGCTGATGGAGAATGGCCCTGTCCAAGCCCTCATG AGAGATACCGCCGGCATGGGACCCACTCAGTCAAGATCACAGGATGGGGAGAGGAGACGCTGCCAGATGGAAG SGGCAAGCGCCAGGCCACTGCCCACTGCCCCAACAGCTATGTTAATAACAATGACATCTACCAGGTCACTCCTG GAGGTGCATGAGGACTTCTTCCTATACAAGGGAGGCATCTACAGCCACACGCCAGTGAGCCTTGGGAGGCCAG GCGCGCGCGTCAATGAGTGCGACATCGAGAGCTTCGTGCTGGGCGTCTGGGGCCGCGTGGGCATGGAGGACAT GCCGCGGTGGCGTCTCGATGGTGCCTGGTTCCTGCGTCGCCGAGGGGTGGTGTCTGACCACTGCTACC GCTGAACCCAGGGGAGGTGCTTCCCACAGCCTTCGAGGCCTCTGAGAAGTGGCCCAACCTGATTCATGAGCCT

SGPR414 SEQID 5

FIG. 1C

GACCAAATTTACATATTGAGATTATCAAACAGTGCCAAGTGATTTTTGAATTTTTTGGCAGCAGAGGGCGACTGAG AAGGAATATAAGCATTTGGAGATTTTTAATCAAGTAGTGTGTGCACTTATTAACTTAGTGATTGCCCAAGTTCAAGT GCTCAGAAAGGAACATACTCTCAAAATATTTACTTACATCAATTCCTGGACACAGAGGCAATGTCTATGCTGCTTC ATGTGCGAGAACTGCGCAGACCTGGTGGAGGTGTTAAATGAAATATCAGATGTAGAAGGTGGTGATGGACTGCA ATGAGAAGGAAAAACTCTTACTATGTGGCAAAAATTTTTCAAATTCAGTTTCCCTTATATACTGCTTACAAGCAT IGGATACAACATTATGCTTTGATAAAGAAAGCCTAGATCTTGCATTTAAGTACTTTATGTCACCTACTTTGACTATG CGGACACAGAAACGTCCATTGCAAAAGAACTTGCAGACTGGCTTATTAGCAACAATGTGGGGGGGCATATATTG FACTCAACATATTGACTGTATTTGGGCTGCAGCACAGTTGAAACATTGTAGTCGGTATATACATGACTTATTTCCTT SCTCCGGGACCAGCTTTGTAAACATTGTACTACCATTAACATAGATTCCACGTGGCAAGATGAGAGATAATCAAGC **AACTCTACAAGAATTTGTAATCTGACTGAGGAGGAATCTTCAAAGAGTTCTGATCCTTTTAGTTTATGGAGTACAG** CTTTGAATATGGAACTCCTGAAACTTTGCCATTTCTTATAGCACATGCGTTTATTACAGTTGTGTCTAATATTAGAA CTCGGATCAGGAGTTACGACAGAGTGCAGCTCGTAACATGGCTGACTTAATGTGGAGCACAGTCAAAGAACCAT 4GGTTGGCTGGATTGAGTCAGATAACAAATCAACTCCATACCTTCAATGATGTGTGCAATAATGAATCATTAGTAT <u> ATGGCTACATATTCCCGCTGTCATGCAGCACATTATACCTTTTAGGACCTATGTTATTAGGTATTTATGCAAGCT</u>

GAGCAAGTTGACATCTTATGGCATTGTTTAGTAGAAGATTCTGAATGTTATGATGATGCACTCCATTGGTTTTTAAA GCCTTGAAAACTTGGGAAACAACAGATCAGTAGTAATTTCACTTCGTCTTCCAAAACTATTTGGTACTTTTCAG GCTAAAACCTGAAACAATTAGCATGACTGGCTTAAACCTGTTTCAGCATCTCTGTAACTTGGCTCGATTGGCTACC AGCAAGAATTTATTAGTAAGTGCATGGAGAGTCTTATGATAGCTTCTAGCAGTCTTGAACAGGAATCACACTCAAG SCTAAGGGTTGTATGCCAGCCAGCTGGACTTCCTGACAAGATGACTATTGAAATGTATCCTAGTGACCAGGTAGC <u> AATCATTGCCATCAGTAGATAATCGAATGCGGATGCTGGATGCTTGTTCACACTCTGAAGACCCAGAACATGATAT</u> TCTGGTGATGTCAGTCGAGCAGCTATCCAGTATATTAACTCCTATTATATTAATGGTAAAACAGGTTTGGAGAAGG CAGTAGTGGGCATAGTGATGGATCTAGCAATGAGGTTAATTCTAGCCACGCAAGCCAGTCAGCTGGGAGCCTG CCATCACCACCACCACGATGGGCATATGGTTGATGTATGCTAAGTGCAGATGATGTCAGTTGTAGTAGCTCCCA CTTAGTCAAGGACCTGTAGTTCATAAACATCAATTCAACAGTAATGCTGTTACAGACATTAATTTGGATAATGTTTG ICTGAGACAGTGGCAAATTGAAGGCACTGGTATTAGTAGTCATTTGAAAGCACTGAGTGACAAACAGTCTCTGCC CACTCATCAAGAATTTGGATCCCGTACCACTTAGACATCTACTTAATCTGGTCTCAGCTCTTGAGCCAAGTGTTCA CATAATCCTCCCAAAAGCAGTTGTGGTACAGATCTTCGGAATAGAAAGTTAGAGAGTCAAGCAGGCATTTGCCTG CAGTITIGGGAGCAGTIACGATACACACTGGATAACAATGTGGGCAGAAAAAAAAGAACTGAACATGATGAAGCTTTTC **AAGTTCAAGTTCGTCTTCAATTCTTGACTTGTGTATTTTCAACTCTGGGATCACCTGATCATTTCAGGTTAAGTTTA** TTCAGGGGAAATGAATGCTACTCATATAGCACAAGGGTCTCAGGAGTCTTGTATCACACGAACTGGGGACTTCCT GGTTAGTGCAAAATCAGAAAAAAAAAATATGGCTGATTTTGATGGTGAAGAATCTGGATGTGAAGAGGGGGGCTAGTTCA TCAAGTTCGAAGTAAAGATCAACATGCTATGGGTATGGAAACCTACAAACATCTTTTCCTGGAGAAGATGCCCCA CTCATGGTTATAGAAAGAGGACTCCTTATGCTGAAGACACATCTGGAAGCGTTTAGGAGAAGGTTTGCATATCA TACTGAACAGACACTGTACTTGGCATCCATGTTAATTAAAGCACTGTGGAATAACGCACTAGCAGCTAAGGCTCA TTAGAAGAACAGCTCCATCACCTTGGTCACCTGCAGCTAGTCCTCAAAGCAGTGATAATAGCGATACACATCAAA AGTGCCTATGATGGTTGTTCAAATTCTGAGCTGTGTGGTATGGACCAATTTTGGGGCATTGCTTTAAGAGCACAA GTGGAGGTAGTGACATTGAAATGGATGAGCAACTTATTAATAGAACCAAACATGTGCAACAACGACTTTCAGACA CAGAGGAATCCATGCAGGGAAGTTCTGACGAAACTGCCAACAGTGGTGAAGATGGAAGCAGTGGTCCTGGTAG GCAGTGAGGTACAGTCAGAAGACATTGCAGATATTGAAGCCCTCAAAGAGGAAGATGAAGACGATGATCATGGT TGGGGAGACTATTGGGAATGAATTATTTAATTGTCGACAATTTATTGGTCCACAGCATCACCACCACCACCACCACCACCA SATTAATTCACATGCGGAACTGACATCTCACCTCCAACAACATCTTCCCAATTTAGCTTCCATTTACCATGAACAT CAAGAAAGGAAATACTTTGTTGTGGGATATAGTCCAAGATGAAGATGCAGTTAATCTTTCTGAAGGATTAATAAT GAAGCAGAGAAACTTCTTTGTTCGTTAGTATGTTGGTTTACAGATAGACAAATTCGAATGAGATTCATTGAAGGTT

TTATCCCTGTCAGGGCTGGATGGAGGAGACTCAATCATCGTTCTTTTCTGCTATTGGCTGCCTCAACATTATTGA CAATGTGGCTCTTGGTGAGTTGGGCTCATTGCTGTTCTTTAGTGAAATCTAGCCTTGCTGATAGCGATCATTACA 4GATTGGCTAAAGAAATTGACTCTCCTTATTCCTGAGACTGCAGTTCGTCATGAATCATGCAGTGGTCTCTATAAG 4ATGTAGAAGATGATGGGCTTACAGGACTCCTAAGGCTTGCAACAAGTGTTGTTAAACACACAAACCCCTTTAAAT TTCAAGGGAAGGACAGGAATTTTTGAGAGATATCTTCAATCTCCTGTTTTTGTTGCCAAGTCTAAAGGACCGACA <u> CTACTATTCAGCAACTTTATATGATACCTGAGGCAAGACAGGCTGTCTTCACTGCCAAGTATTCAGAGGATATGA</u> 4GCACAAGACCACTCTTCTGGAGCTTCAGAAATGTTTACATATTTAATGGAGAGTGAATGCAAAGCATATAATCC **AATTTCTTCCTGATGCTCAAGCACTCAAACCTATTAGGATAGATGATTATGAGGAAGAACCAATATTAAAACCAGG** CTCGACTTAGATGCCTTGGCAAGACATTTGGCTGACTGTATTCGAAGTAGGGAGATCCTTGATCATCAGGATGGT STGGCCTCATGAAGATGTCCGTGCTGAATGTAGATTTGTTGGCCTTACTAACCTTGGAGCTACTTGTTACTTAGCT SCCAAGTGGCTGATATGAAGAACATTTATGAATCTCTTGATGAAGTTACTATAAAAGACACTTTGGAAGGTGATAA SATGTATACTTGTTCTCAATGTGGGAAGAAGTACGAGCTGAAAAAAAGGGCATGTTTTAAGAAATTGCCTCGCATI 4CACGAGTTAACAACAGATTATGATGAAAAAGCACTTCATGAGCTTGGTTTTAAGGATATGCAGATGGTATTGTA SATGCAGAAAATCTGTCTAGGCGGGTCTGGGAGCTACTGATGCTTCTTCCTACATGTCCTAATATGTTGATGGCA **ATGTAAAGAGTATITITIGGTIGTTATGCAAATTAGTTGACAACATACATATAAAGGACGCTAGTCAGACAACGCTC** 'GAGAACTACAGGCTAATACACACAGGGTTATGGCACAACACATGCAGTCCCATGCACGTTATAAATGGGATTA 4GAAGTCAGTGATCATTCAAAAGACTCAGAGAGCTATGAATATGACTTGATAGGAGTGACTGTTCACACAGGAAC CTITITAATGATGCTGAGGTAAAACCTTTTGATTCTGCTCAACTTGCATCTGAATGTTTTGGTGGAGAGATGATGACGA GAGTTTGGTCAAAGCAACCGAAAAGGAGAGTTTCCTGGAGGCCTCATGGGACCTGTCAGGATGATTTCATCTGG ACAGCCAAAGTGCAAATCACATTCTTCAAGAGCTGCCGCTTACGATTTGTTAGTAGAGATGGTAAAAGGGGTCTGT TNAGTITCAATACTATGAGATACACATTTAATATGGTCACGATGATGAAAGAGAAAGTGAATACACACTTTTCCTT SCAAGACCTATGATTCTGTTACAGATAAATTTATGGACTTCTCTTTTGAAAAGACACACAGGGCATATATGCTGTTT **3GCAGATGGTGGACACTATTATAGCTTTATCAGAGATATAGTAAATCCCCATGCTTATAAAAACAATAAATGGTAT** ICTTTGGGTGCACCAAGGAGAGAGGGGAAAGGGGGAAGGTGTTCAGCTGCCAGCATCTTGCCTCCCACCCCTC TTCCAGAATATCTCAGATGAGCAGAGTTTTAAAGCTCAGTCTGATCACAGGTCTAGACATGAAGTTTCACATTATT STAATTACAAACAATGTTGTATCCTTGGATTGTGAACATGTTAGTCAAACTGCTGAAGAGTTTTATACTGTGAGGT -AGACCTTTCTGTAAAACATACACCATGGATAAGCAGCCTCTGAATACTGGGGAACAGAAAGATATGACAGAGTT AGAAGGACAACATTCCAATGCTTTTGCTTTTACAAGAGCCTCATTTAACTACTCTTTTTGATTTATTAGAGATGCTT 3CATCATTTAAACCACCCTCAGGAAAAGTGGCAGTGGATGATAGTGAGAGCTTACGATGTGAAGAACTTCATCT TTTACTGATCTAATTACCAAAATCGAAGAAATGTCTCCCGAACTGAAAAATACCGTCAAAAGTTTATTTGGAGGT

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GACTGACATGGCAGCATTAACAGGAGGAAAGGGATTTCCCTTCTTGTTTCAACATATTCGTGATGGCATCAATATA SATTGATTCTAATGACAGAGTCTTTCAACACTTTGCACATGATGTATCACGAAGCTACAGCTTGCCATGTGACTGG 4GATTTAGTAGAACTTCTGTCAATATTTCTTTCGGTTTTGAAGTCTACACGCCCTTATCTTCAGAGAAAGATGTGA TCTCAGTGTCTAGATTGGTTGGCAGTGCAGACACCCCGAAATAAACTGGCACACAGGCTGGGTCTTACAGAATATG **AATACCCTGGAGCAGTAGAAGAACTGTTTAACCTGATGCAGCTGTTTATAGCTCAGAGGCCAGATATGAGAGAAG** TACAAACGCATGGAACCAGAGGAAGAAAATGGCAGAGAATACAAATTTGATGTTTCGTCAGAGTTACTAGAGTGG GTCCTAGAGACATTTATTCATTCTAAAGAAAAGCCCACGATGCTTCAGTGGATTGAACTGTTGACGAAACAGTTTA TTAAGTGCCCTAATCAAATTGTGAGACAGATGTTTCAGCGTTTGTGTATCCATGTGATTCAGAGGCTGAGACCTGT GCATGCTCATCTCTATTTGCAGCCAGGAATGGAAGATGGGTCAGATGATATGGATACCTCAGTAGAAGATATTGG TCCGCCTTATTGTTCAGAACCCAGTGGTAACCAAGAACATTGCCTTCAATTACATCCTTGCTGACCATGATGATCA CTGCATTCACACGACAACTGGCTTCTCACCAGAACATCCAGTGGGCCTTTAAGAATCTTACACCACATGCCAGGC ATAATAGTCAGGCAGCTTGTGAGTGGTTTTTAGATCGTATGGCTGATGACGACTGGTGGCCAATGCAGATACTAA GCAAGCTATATCTACAATGGTACATTTTTACATGGGAACAAAAGGACCTGAAAATCCTCAAGTTGAAGTGTTATCA CCACTCAGTCCAGACACAACAGTAGTCCTACATCAGGTCTACAACGTGCTCCTTGGTTTGCTCTCAAGAGCCAAA GCAATAGCTACAAATCACAATAAACAGGCTTTGCTTTCATTTTGGTACAATGTCTGTGCTGACTGTCCAGAGAATA CCCTTGAAAAGATGATAGCTTTTAGTTGCTCTTTTGGTTGAACAGTCTCGATCAGAAAGGCATTTGACATTATCACA GGTGGACCTCCAGGAATGCCTCCCTTTGCATCTTATATTCTGCAGAGGATATGGGAGGTGATTGAATACAATCCT CTITATGTTGATGCTGCTGTTCATGGCACTACAAAGCTAGTGCCCTATTTTAGCTTTATGACTTACTGTTTAATTTC CAAAACTGAGAAGCTGATGTTTTCCACATATTTCATGGATTTGTGGAACCTTTTCCAGCCTAAACTTTCTGAGCCA GGATGTGGTGCTTTTTAACCGTGGGATGCTGCCAGCGTACTATGGCATTCTGAGGCTCTGCTGTGAGCAGTCTC <u> AGACAAACTTGTAATCTGATTTTCAGCCTGTGTCGATACAATAATCGACTTGCAGAACATATTGTATCTATGCTTTT</u> GAAAACTGGGTCGAGCGGTTTCTTTTGGCTCACAATTATCCTAGAGTGAGGACTTCTGCAGCTTATCTTGGTG **AACAAGCATTAATCCAGTGGCAGGAGCGAATTGAATTTGCCCATAAACTGTTAACTCTTCTTAATTCCTATAGTCC** <u> STGTAGTTGTATTCCCAGTACATTACCAGATCCTAAAGCTGTGTCCTTAATGACAGCAAAGTTAAGCACTTCCTT</u> TCCCTTATACCAAGCAATTCATTCCGTCAGATGTTCCGGTCAACAAGGTCTTTGCACATCCCAACCCGTGACCTT

GCTTTGCTCCTGTCAGTACACACTCCCAAACAGTTAAACCCAGCTCTAATTCCAACTCTGCAAGAGCTTTTAAGCA CTCAGTCTGCTATGTCAAAAAACTGCATCAAGCTTTTGTGTGAAGATCCTGTTTTCGCAGAATATATAAATGTATC AATGCAGGACTTGTCTGCAACAGAGAAACTCACTCCAAGAGGCAAGAAGCAAAGAAGAAAAAAACTAAAGATGATG CTGGTTCCCTTTCTACAACACAACCATTGTACTTACCATCACAGTAATATACCAATGTCTCTTGGACCTTATTTCCC CTAATGGATGAAAGAACTTTTTAAACAACAACATTGTCTACACGTTCATGACACATTTCCTTCTAAAGGTTCAAAG <u> AAGGAGCAACTCCCATTAAAAGGCGGCGTGTTAGCAGTGATGAGGAGCACCTGTAGACAGCTGCATCAGTGAC</u> TATCTCTGCAGTTCTGTCTGACTTAGCTGACTTGAGAAGCTGTGATGGCCAAGCTTTGCCCTCCCAGGACCTGA TTATCATCAGTTCATCCATCTATTATGCCGAGTTGCAATCAACTGTGAAAAATTTACTGAAACATTAGTTAAGCTGA GTAAAGAATTTAAAGACCTCCACTGTTCCAAGGATTCTACCCTAGCTGAGGAAGAATCTGAGTTCCCTTCTACTTC <u> SGTTGCTTTATCTCTCAGTTGTGGCCATTCCAGAGGACTCTTTAGTCATATGCAGCAACATGACATTTTAGATACC</u> TGTCGAGAAAATATCAAGCTAATAGGAGGGAAAAGCAATATTCGGCCTCCGCGCCCTGAACTCAATATGTGCCT ICAAGTGTTTTCTGAAGCAAACTGTGCCAATTTGATCAGCACTCTTATTACAAACTTGATAAGCCAGTATCAGAAC CTACAGTCTGATTTCTCCAACCGAGTTGAAATTTCCAAAGCAAGTGCTTCTTTAAATGGGGACCTGAGGGCACTC ATGAAAACAGAAACCAGGGAGGTCCTGACCCCAACGAGCACTTCTGACAATGAGACCAGAGACTCCTCAATTATT GATCCAGGAACTGAGCAAGATCTTCCCTTCCCTGAAATAGTTCTGTTAAAGAATACCGAATGGAAGTTCCATCT CTGTGTAGGACCATTGAATCTACAATCCATGTCGTCACAAGGATATCTGGCAAAGGAAACCAAGCTGCTTCTTGA STTGCCCACAATGGTGGAAACCAGTAAGGGCAAAGATGACGTTTATGATCGTATGCTGCTAGACAACTACTTTTC GTGTCCTAGTTGCCTATGAAGGTTTGCCACTTCATCTTGCACTGTTCCCCCAAACTTTGGACTGAGCTATGCCAGA CGTTTTCAGAAGACATGTCAAATATCAGGTCACAGCATGCAGAAGAACAGTCCAACAATGGTAGATATGACGATT ICCAGAACTTAGAAATGCCTGTATAGATGTCCTCAAGGAACTTGTACTTTTGAGTCCCCATGATTTTCTTCATACT

CATCGAACAACAAGGCCATGACAGATCCCTCCAGAAAGTATTTAACCAGCAGTAGAGAAAAGCAGCTGAGTTTGA ITTGGAAACTAAAGATGATATTCCATTTCGAAAAGTTCTTGGTAATCCGGGTAGAGGATCGATTAAGACTGTAGCA ATGTCTCCTCTGAAGATACATGGTCCTATCAGAATTCGAAGTATGCAGACTGGGATTACAAAGTGGAAAGAAGGA 4GCAGGACCTCACAGAAGGAAACCAGCAGGCAGCTTTCTTACTCAGACAATCAGGCTTCTGCAAAAAGAGGAAG rcctttgaaattgtagaaaaagagaataaagtcagcctagttgattcactacaatactggaggaattccaaggatat TTCAGCTAAGTCATAACATTAAAAATGTGGTGCTTCGACCCAGTGGAGCGAAACAAAGACGCGCCTAATGTTAACTC GCAAGATAACAGCTTCTTGTCTATTGACAAAGTACCAAGTAAGGATGCAGAGGAAATGAGGTTGTTTCTAGATG CAGTCCATCAAAACAGACTTCCTGCAGCCATGAAACCGTCTCAGGGGTCTGGTAGTTTTGGAGCCATTCTGGGC SGAAGTGGAATAGCTCGGACGATTCCTTTGACATCTACTTCAACACCTCTTAGATCAGGGTTGCTAGAAAT >SGPR430 SEQID 6

SACCAAAAAGGATTTACTCAAGAAGGTTAAAAATGCCATTTCAGCTACAGCAGAGAGATTCTCTGGTTATATGCAG 4TGAGAATAAAGAAAACAAAACTCCAGAAGGATCTCAGGGAGAAGTTGATTGGCTCCAGCAGTATGATATGGAGC GGGCAGCAAGTCATCCAAGATACCTGACCCTGTCATCTCATTGCACTGAAAATACAAAACCACCTTTTACCC SCGAAGAGAGCTTCTAGCAGCTGTCTTGGAGATAAGTAAGAAGAGATGCTTCACCATCTCTGAGTCATGAAGATG &AAATCCAAGAGGCTGCCGTGCAGAGTGATCGAGATCGGAGTGGCTACATCTTCTTTATATGCACAAGGAGATC GAAGTTTGGGATTTCTTCCTCAGCCAGTTCCTCTTTCTGTTAAAAAACTGAGGTGTAACCAGGATTACACTGGCTG 4GACTGAACCTGTTTCTGGAGAAAAATTCACCAGATATTTCAGCTACCAGAGCATACACTTGCCCTGTTATTAC CTTCTACACCTTCAAAGAAATTCACCTTCAAATCCAAGAGCTCCTTGGCTTTATGCCTTGATTCAGACAGTGAGGA ITCTGATGAGGACTCTGGAAATGAGGATGTTTTGATATGGAGTACACAGAAGCTGAAGCTGAGGAACTGAAAAG SAATAAACCAAGAGTGCCCCTTTCCTCTCACCAACAGCAGCAACTGCAGGGCTTCTCCAATTTGGGAAATACCTG CTTTAGGGCCGAAGAACTGGAGTATTCTTGTGAGAAGTGTGGTGGGAAGTGTGCTCTTGTCAGGCACAAATTTAA TGGTTGGAGTGCACATATGGCAATGTCTAGACCATTGAAAGCCTCTCAAATGGTGAATTCCTGCATCACCAGCC **ATGATAAGCCAACTAGCAGCCCAGATACCGGATTTGCAGAAGATGATATTCAAGAAATGCCAGAAAATCCAGACA** CTATGGAAACTGAGAAGCCCAAAACAATCACAGAGCTGGATCCTGCCAGTTTTACTGAGATAACTAAAGACTGTG TTGATGAGCTGCTGGAAACAGAAAAGAACTCTCAGTCACTTAGCACGGAAGTGGGGAAGACTACCCGTCAGGC GAGCTAAAACGTTCTGTGGCCCTCAGCCAGAGACTTTGTGAAATGTTAGGCAACGAACAGCAGCAGGAAGACC GTGAAAGGGAAGAGCAAGAGCTTCAGCAGGCACTGGCTCAGAGCCTTCAAGAGCAAGAGGCTTGGGAACAGAA GATCCAAGGAACACTCTTCTGGTGGCACTAACTTAGACAGGACTAATGTTTCAAGCCAGACTCCCTCTGCCAAAA TAATTTGGAGTTTGAGGTTCAGCACTCCATCATTTGTAAAGCATGTGGAGAGATTATCCCCCAAAAGAGAACAGTTT 4GAAGATGATGACCTCAAAAGAGCTACCGAGTTAAGTCTTCAAGAGTTTAACAACTCCTTTGTGGATGCATTGGG AACAGTCAGAAGAGAATAGGACATCAGGTGGGCTTTTACCTTTACAGTCATCATCCTTTTATGGTAGCAGAGCTG <u> AATGACCTCTCTATTGACCTTCCTCGTAGGAAAAAACCACTCCCTCGTTCAATTCAAGATTCTCTTGATCTTT</u> CTATATGAATGCTATTCTACAATCTCTATTTTCACTCCAGTCATTTGCAAATGACTTGCTTAAACAAGGTATCCCAT rggaaaaaattcaaaattatgcccaatagagcctgacaagtctgaattggaaaactcaggatttgacagaatga CAGGCTTCCTAGGGTCCTCATTCTCCATTTGAAACGATATAGCTTCAATGTGGCTCTCTCGCTTAACAATAAGAT

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ACTECATCATAGACCAAATCTTCACAGGTGGCCTGCAGTCTGATGTCACCTGTCAAGCCTGCCATGGCGTCTCCA <u> AGGAACTGCGGGTTATCTACCAGTGCTTCGTGTGTGTGGAACCCCAGAGACCAGGAAAAGCAAGGCAAGGCAAGTCC</u> CCCAACCTGGGAAACAACCAAACCAGAATTAGAACTGCTGGGGCACAACCCGAGGAGAAGAAGAATCACCTCCA STECACCTEGTGTGGATACATGCCCGCCATTTAGCAGGGTACAGGCAACAGGATGCCCACGAGTTCCTCATTGC FCCTTCAGTGAGTGTGCTAGTTGCAGTTCCTCTGGTTGCAGGCCACAAAGGGCCAGGCATTTATTGAAAGGATGCT GGGGAAGGCGGGGGGGGGGGGGGGGGGGGGGGGGGGAAGGTGGACGCCACCGAGAAGGTGGACG COGATOCOTOTOCOGOTOGOAGTOGTOGOGOCOGOCOGOCOCOGOCOCOGOCOCOAGACOCOGOTOGTA TGGTCATCCACGTGAGTGTGAGTCACCAGGATTGATGACAAAAATAGAAGAGGGATTGGAAGACAGTGAGCCAG 36ATGAATTACCTTTTGGTGTAATTGGTTCTAATATTGGTGATGCACATTTTCAAGAATTCAGGGCTGGAATCTCC FGGAAGCCTGTGGTAGATCCTGATGACCCCATTCCTCAGTTCCCTGATTGCTGCAGCAGCAGCAGCAGCAGGAGT SCGGGGAAGGTGGACGCCGCTGGGAAGGTGGAGACGGCGGGGGGTCCGGGCCGCCGCCGGGTTGAGCT 36AGCCCGAACCCGAGCCGGTCCGGGAGGCGGAGCAGGAGCCGAAGCAGGAGCTGGAGGATGAGAACCCAG SATGTGGATTTTGCCGTGGGCCACCAGGCTGTTCTCACGTGAACAGCTTTAAGGTGGGAGAGAACTGGAGGC GCTTTACGATCGGTTTAAGAGGACTCATCAATCTTGGCAACACGTGCTTTATGAACTGCATTGTCCAGGCCCTCA CTGGTCTGTGAGATGTCGTCGCTGTTTCGGGAGTTGTATTCTGGAAACCCGTCTCCTCATGTGCCCTATAAGTTA STGTGAGTCCCAGGACCTTACAAATATCAGTTTTTGTGTTATCAGAGAAATACGAGGGTATTGTTAAATTTGAATC GTGCCAAACTCTTCAATAATGATGGAGTCTGTTGTTGCCTGCAAAAACGGGGGCCAGTGAACATTACATCAGTG1 GCATCTGCCATGTGTGTGGCACCCATCTGAACAGACTCCACTCTTGCCTTTCCTGTGTCTTCTTTGGCTGCTTC CGGAGGAGGTAGAGGCGGGGGGGGGGGGGAGAGGCGGAGGAGGAGGTGGAGGCGGAGGCGAAGGTGGA TTGAAATTACAAGCCTCCACCTCAACAGAGGTTTCTCACCAGCAGTGTTCAGTGCCAGGCCTTGGTGAGAAATI SCCACACGCCGATACTGAGAGATTTCTTTCTCTGACAGGCACCGATGTGAGATGCCGAGTCCCGAGTTGTGT <u>ATGACACTACTTGCTCCCTGGTACACAGGCCCCATGATCCCCATGGATGTTAATGAGCCCAGCTCCGTGACCAC</u> 4CGGAGAAACACATTCACGAGCACGCAGAGACGAAACAACACACTTAGCAGTAGACCTGTATTACGGAGGTAT <u>GGCTCCTACCCTCAGCTCTAGCCTGCAGCATATCTCCTCATTCCTGGCCACTGGTAAGAAACTTTCCCTCCATT</u>

GGGAGGGAGAGCAGTGTGAACGGGGAAAGCCACATACCAGGAATCACCACCCTCACGGACTGCTTGCGGAGGT GGCGCAAGATCACTACATACATTTCCTTTCCTCTGGAGCTGGATATGACGCCGTTTATGGCCTCAAGTAAAGAGA GCAGAATGAATGGACAATTGCAGCTGCCAACCAATAGTGGAAACAACGAAAATAAGTATTCCTTGTTTGCTGTGG TTAATCACCAAGGAACCTTGGAGAGTGGCCACTATACCAGCTTCATCCGGCACCACAAGGACCAGTGGTTCAAG 1GTGATGATGCCGTCATCACTAAGGCCAGTATTAAGGACGTACTGGACAGTGAAGGGTATTTACTGTTCTATCAC CCACGATAGACCCATGCTGGGACATTAGTTTGGACTTGCCTGGCTCTTGCACCTCCTTCTGGCCCATGAGCCCA AAACAGCTCACAATGAATAAATTACCTGTCGTTGCCTGTTTTCATTTCAAACGGTTTGAACATTCAGCGAAACAGA ITACGAGGCCAGAGCACTTAGGAAGCAGTGCCAAAATCAAATGTGGGTAGTTGCCAAAGCTACCAGGAATCTACC 

SGPR495 SEQID 8

AAATACAGAAGGAGGAAAATGCAGAAATTTATCTGTAAGAGGAATTACAAATTTAGGAAATACTTGCTTTTTTAAT GCTTTGTCAGAAGGCACCTCGATTTAAAGATTTCCAGCAACAGGACAGTCAGGAGCTTCTTCATTATCTTCTGGAT GAAGACTCTTCAGATGATATTGCTGTAGGTTTAACTTGCCAACATGTAAGTCATGCTATCAGCGTGAATCATGTAA **AAGAGTCAACTAATTCATGACCGAAAATGTATTAGAAAATTGTCATCTGGAGAAACTGTCACATACCAGAAAATG** TCAAGATTTTTCCTTCCTCAGACTCTCAGCTGGÄCCCATTGGTGGTGGAACTTTCAAGGCCTGGACCACTGACCT CAGCCTTGTTCCTGTTTCTTCACAGCATGAAGGAGCTGAAAAAGGACCACTTTCTCCTAAAGTTCTTTTAATCA FAGTACTTACTTCTGATATTTGGTTGTGCCTCAAGTGTGGCTTCCAGGGATGTGGTAAAAACTCAGAAAGCCAAC **AAACATGCTTCTAAAACAACAAGTGCATTTTCTAGAATCATGAAACTTTGTGAAGAAAAATGTGAAACAGATG** GCAGTGAGGACAGAAGAAACAAAGCGAATACAAGCTAGCATTCTAAAAGCATTTAACAACCCAACTACTAAAACT STGTTATGAATGTGATGAAAAATTATCAACGCATTGTAATAAGAAGGTTTTGGCTCAGATAGTTGATTTTCTCCAG CGATACAGTGGCAATGTTACTATAGAAAATATTCATCAACCTAGAGCTGCCAAGAAGCATTCTTCATCTAAAGAT 

'ACTGATGACAGTGAAAAAGAAGCCAGCCATTCTGAAAGCAATGTTGATGCTGACAGTGAGGCCTTCAGAATCTGA

**AAGTECTTCAAAGCAGACTGGGCTGTTCAGATCCAGTAGTGGATCCGGTGTGCAGCCAGATGGACCCCTTTACC** 

CTCTGTCAGCAGGTAAACTGCTGTACACCAAGGAGACTGACAGTGGTGATAAGGAAATGGCAGAAGCTATTTCT

AGTGACACTTACTTACAGGTGGTTCCAGAATCAAGAGCACTTAGTGCACAAGCCTACCTTTCTATGAAAGAG SAACTTCGTTTGAGCAGCACTGTAACTGGGGATCAAGATTTTGACAGAGAAAATCAGCCACTAAATATTTCAAATA 4GAAGGAGTTTATACTAATGCCAGGAAGCAATTGCTCATTTCTGCTGTTCCAGCTGTCCTAATTCTCCACCTGAAA 'ATAACTACTTCTAAAGAATGTTCAATTCAGTCCTGTCTACCAGTTTACATCTATGGAATTACTAATGGGAATA <u> AGATTTCATCAGGCTGGCTTGAGTCTTCGTAAAGTAAACAGACATGTAGATTTTCCACTTATGCTCGATTTAGCAC</u> ATTIATGITITITAGAGGGAAGCATTTGAGGTCTTATAGTCCCCAAAATGCTTTTCAGACCCTTTCTCAGAGCTA CATTCTGCTCTGCTACTTGTAAGAATGCAAGTGTGGGAGATAAAGTTCTCTACGGTCTCTATGGCATAGTGGAAC CATAACACTAAAAAGAAAAATGTGCCTGGTTTGAAAGCGGCTGATAGTGAATCAGCAGGCCAGTGGGTCCATGT **4TAGTGGCTCGATGAGAGGCCACTACACTGCTTATGTGAAAGTGAGAACACCCTCCAGGAAATTATCGGAA** 

#### SGPR407 SEQID 9

FIG. 1K

TGGCTGGTGTTGGTGCCTCTGAAACTGACAGGAAGTCCCCACAGATGGAGGGCCCAGGAAGAGAGGGCCTGGCCA GAACCACTCAGGCAGCATCCACTGTGGCCACTACACAACCCTGTGCCAGTGCCAGACTGGTTGGCACGTTTACA <u> ATGCCCTTTGTGATACCGAGCAAAACCTTGCCATGGGACCCACTGGAACTCAAGATTTGTTATCAGCAAAATCGC</u> 7ATGGGAGCGGCTACACCTTGAAATCAACCACGGAGGCCGCGGGGCTCCACCAGTCCCTGCCCATGGTCCAGC GCTGCAGCTGGTGTCTCCAAAGGGTCACCATGAGGCGGGTTATGGGTGTGCAGGACAAAGCTGGAAACAGGAA ATGGAGTATCCAGTCCCATACTTTAGATCCCCGAACAGGACTCTGATCCCAGAGAGAATTTGGTCAAACCCATTA ATGACTCTTGTGTCTCCCCTAAACACGCTGCGGGACACAGAAGGAATAGAACTCACAGTTATGAAGGCTCTAGTT SCTTGCAGACGGCCACCTTCTCACTCTGTCTTCACACAGTCCTTCTGGGCATGTCTGGATCCTGATCTTCT \*\*CCTCTCCACCCACCAAGGGGAGTGCTCTGCTAAAAGAGTCTGAGTTAAATGATGATGCTGACTGGGCCAACCTA ATGTGGAAGCGTTATCTGGAAGAACAAGAGGACAGCAAGATGGTGGATCTGTTGTGGGCCAGATGAAAAGTTA AGAGTTAGAGAATGCCTCAGGGACTTTGCCAGTGACAAAGTCGGAAGTCCTGTCTACCAGCTGTGTGCCTTTG AGTTACCAGAACCTGGAGGTTTGGAAAAGAAACATGAAGAGCTGAGACTCAGACCTCTGAAGGAGGAGTACCAT CCATATCCCTCCCCCGACCCATCAAACTTTCCTACCTTCTTACGCTGTCTGAATGCTTTCTCTGCAGCTGTCTTCT CCCAAGAAAGGATTTGCTGGGGGCAAGGTGTCTCTGCGGGATTGTTTAAGCCTTTTCACCAAGGAAGAAGAGCT CCAGATGCTGCTGCTGGGGCAAAGACCTGTGATAGGTGATACAGTCAGCAACAGCCAGACAACTAGGGACAAG ATCTCCCACAGCCCTCATGGCATAAGCCCGAGGGCTTAAAGCCAGCAGGATACCCAAGAGTTCCTGACATTCCT

AGTGGGATGGAGCCACAGAAGTCTGCACCATTTGCAGCAGGGAAGGGTCTGGCCCCTCCTCTTCCTGTGTGCAA ATGCCCCAGAGTTTTTACTGGGGTTTTCCTGCTGGAGGGAACTTGTCTGCTTTAGAAATGCTGCCTGATGGACCA GCACCAAGGACGTITCTICAGAAGAAAAGCTGTCTCTTTCCCCTGTTCTTTACATTCTTTTGCATAAGGCAGGTA <u> AACTCTTCCAGCCTGATGCTCATGGATTTCTAGTGAAGAAAGTTCATGCTCCAACAAGGGGGCATCGTGTTTATCAT</u> GGAACCAAGACAGCTGGGTGGGAAGGGCTCCCTGTCAAAACTCCAACCAGCCTGTGCACTGGGAGGAATGAAC CCAAGGACAAATTCATTCCCCAAGGACAAATTTGTCCCCAAGGACAAATTGAAGGTGATATTGTCCCTGCTGACA CCTGAGATTCAAACTACGAGTTTACAAATTTGAGGAAGAGCTTTGGTCCAGGGCAGGCTTGGGGAAGAAGTG ACAACCACTCATCTAGGCAGATGCCCTGGGGTGCCGCTGGGGTGGCATGCCAGCATCCATGTAAACTGCCAG TCCCTGTCTCAGTTTTCCTCAATGATTCTCAACCAGAGGAAGCAATACCTCCTCAATCCCTGCTCCCGGGTTCC AATTGTTGCAGAGTTGACACCTCCAAAATTGTCATTTGGTTTCCTGAACACAGGTTCAGAGTTCAGTACTTCCTACT ATGTATGAACTAGACCGATTATTT

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FIG.

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GTCATCAGGACTAAGTGGTGGAGCATCAAAAGGTAGAAAGATGGAACTTATTCAGCCAAAGGAGCCAACTTCACA 'ACATTAAGTGCCATCAAAAGTCAAAATTATCACTGCACAACTCGTAGTGGGAGGTTTTTACGGTCCATGGGTACA <u> AAGAAGAACCATTTCAGGAGAAAATAGTAGTAAAAAGAGAAGTAAAGAAAAGACGGCAGGAATTGGAGTATCAAG</u> FAAAGCTTGATCTGAACCAATGGCTGGCTATGACTGCTAGCGAGAAGACAAGATCTTGTAAGCATCCACCAGTCA GTACATTICICITITGTCATGAATTGCATACTTTGTTCCAAGTCATGTGGTCTGGAAAGTGGGCGTTGGTCTCACCA GGCACAGGAGAAGGATACTAATGGGTAAAATCTTTCGAACATGGTTTGAACAATCACCCATTGGAAGAAAAAGC SAAGATATATTGAAGAGCATGCACTCAAGCACTTTCAAGAAAGCAGTCATCCTGTTGCATTGGAGGTGAATGAGA TGTACGTTTTTGTTACCTTTGTGATGATTATGTTCTGAATGATAACGCAACTGGAGACCTGAAGTTACTACGACG AAATAGTTTCTGTTCAGGTGCCAGCACAAACGCCAGCATCACCAGCAAAAGATAAAGTACTCTACCTCAGAAA SAAATTTGGGAAATACTTGCTATATGAATTCTGTTCTTCAGGTGTTGAGTCATTTACTTATTTTCGACAATGTTTT GAAATGGCACTGTGTGGACTGCAACACGACCGAGTCCATTTGGGCTTGCCTTAGCTGCTCCCATGTTGCCTGT ATGCTAGCAATGGATACGTGCAAACATGTTGGGCAGCTGCAGCTTGCTCAAGACCATTCCAGCCTCAACCTCA ATGAAATATCTCAAAAAGTCAGTGACTCCTCAGTTAAACGAAGGCCAATAGTAACTCCTGGTGTAACAGGATTGA GGTGATGATTCTTATTTCTTACATGACGGTGCCCAATCTCTGCTTCAAAGTGAAGATCAACTGTATACTGCTCTTT ITTGCTATGCTACACTCAGTGTGGAGACTCATTCCTGCCTTTCGTGGTTACGCCCAACAAGACGCTCAGGAATTT CTTTGTGAACTTTTAGATAAAATACAACGTGAATTAGAGACAACTGGTACCAGTTTACCAGCTCTTATCCCCACTT GATTCCAAACTAAGCATGTGCACTATGGATGAAGTATGCAAGGCTCAAGCTTATATATTTTTATACCCAACGAG CTCAAAGGAAACTCATCAAACAAGTTCTGAATGTTGTAAATAACATTTTTCATGGACAACTTCTTAGTCAGGTTACA TACTGAGAATGGACATTCTAAACTTTTGCCTCCAGAGCTCCTGTTGGGGAGCCAACATCCCAATGAAGACGCTG I GT CTT G CAT G T G A CAAAT C C A T A C C A T A G A A C C T T C T G G A C T T G G A G T T T C C A G A A G G T A C GTCAGGACGTAATAACCGAGAGAAGATTGGTGTTCATGTTGGCTTTGAGGAAATCTTAAACATGGAGCCCTATTG CTTTAGAAGGAAAAATCTACGTATGTGACCAGTGTAACTCAAAGCGTAGAAGGTTTTCCTCCAAACCAGTTGTACT STGCAGGGAGACCCTGAAATCCCTCAGACCAGAATGCTTTATCTATGACTTGTCCGCGGTGGTGATGCACCATG AATGCAGTGGAAAAGATATTGCTTCCCAGCCATGTCTGGTTACTGAAATGTTGGCCAAATTTACAGAAACTGAAG 3GAAAGGATTTGGCTCAGGGCACTACACTGCCTACTGCTATAATTCTGAAGGAGGGTTCTGGGTACACTGCAAT ATACCTCGTCTAATGAAATCCTTAGCTGA

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AAATACAGAAGGGAGGAAAATGCAGAAATTTATCTGTAAGAGGAATTACAAATTTAGGAAATACTTGCTTTTTAAT <u> GAAGACTCTTCAGATGATATTGCTGTAGGTTTAACTTGCCAACATGTAAGTCATGCTATCAGCGTGAATCATGTAA</u> 4GAGAGCAATAGCTGAGAATCTGTGGTCAGTTTGCTCAGAATGTTTAGAAGAAGAAGAAGATTCTATGATGGGCAGC TCAAGATTTTTCCTTCCTCAGACTCTCAGCTGGACCCATTGGTGGTGGAACTTTCAAGGCCTGGACCACTGACCT AAACATGCTTCTAAAACACAAACAAGTGCATTTTCTAGAATCATGAAACTTTGTGAAGAAAAATGTGAAACAGATG CAGCCTTGTTCCTGTTTCTTCACAGCATGAAGGAGACTGAAAAAGGACCACTTTCTCCTAAAGGTTCTTTTAATCA <u>ATGCGGGTGAAAGATCCAACTAAAGCTTTACCTGAGAAAGCCAAAAGAAGTAAAAGGCCTACTGTACCTCATGA1</u> GTGTTATGAATGTGAAAAAATTATCAACGCATTGTAATAAGAAGGTTTTGGCTCAGATAGTTGATTTTCTCCAG I A GTA CTT CT G A TATT T G G T T G T G C C T C A G T G C G C G G G A T G T G G T A A A A A C T C A G A A G C C A A C GCTTTGTCAGAAGCGGGTGCATCTACATTTAATATAA

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GAGCAGTTTCCAATCAATGAACACTATTTCGGATTGGTCAATTTTGGAAACACATGCTACTGTAACTCCGTGCTTC <u> ATGACTGTCCGAAACATCGCCTCCATCTGTAATATGGGCACCAATGCCTCTGCTCTGGAAAAAAGACATTGGTCCA</u> TTGCTGACGTGCCTGGCGGACCTTTTCCACAGCATTGCCACACAGAAGAAGAAGGATGGCGTCATCCCACCAAA GAAGTTCATTTCAAGGCTGAGAAAAGAGAATGATCTCTTTGATAACTACATGCAGCAGGATGCTCATGAATTTTTA 

AATTATTTGCTAAACACTATTGCGGACATCCTTCAGGAGGAGAAGAAACAGGAAAAAACAAAATGGAAAATTAAAAA ATGGCAACATGAACGAACCTGCGGAAAATAATAAACCAGAACTCACCTGGGTCCATGAGATTTTCAGGGAACGC TACCAATGAAACTCGATGCTTGAACTGTGAAACTGTTAGTAGCAAAGATGAAGATTTTCTTGACCTTTCTGTTGA **IGTGGAGCAGAATACATCCATTACCCACTGTCTAAGAGACTTCAGCAACACAGAAACACTGTGTAGTGAACAAA** rcctctggaactccggctcttcaacacctccagtgatgcagtgaacctggaccgcatgtatgacttggttgcg GTGGTCGTTCACTGTGGCAGTGGTCCTAATCGTGGGCATTATATCACTATTGTGAAAAAGTCACGGCTTCTGGCTT TGTTTGATGATGACATTGTAGAGAAAATAGATGCTCAAGCTATTGAAGAATTCTATGGCCTGACGTCAGATATAT **ATATTATTGTGAAACATGCTGCAGCAAACAAGAAGCCCAGAAAAAGGATGAGGGTAAAAAAAGCTGCCCATGGTCT** CAAAAAATTCAGAATCTGGATATATTTTATTCTATCAGTCAAGAGAGTAA

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<u> AGATATGAGTACTTTTATTGATGTTGAAGATGAGAAATCTCCTCAGACTGAAAGTTGCACTGACAGTGGAGCAGA</u> SCAAACAGAACAGGCTGATCTTATAAATGAGCTATATCAAGGCAAGCTGAAGGACTACGTGAGATGTCTGGAATG 'GGTTATGAGGGCTGGCGAATCGACACATATCTTGATATCCCATTGGTCATCCGACCTTATGGGTCCAGCCAAGC SAAGCAGAGGATATTATTCTAGTGCTTTCGCAAGTTCCACAAATGCATATATGCTGATCTATAGACTGAAGGATCC 
 IAGTGAGGCTTGGCAGCAGCATGTACAAGAACTATGCAGAGTCATGTTTGATGCTTTGGAACAGAAATGGAA
**GGTACAGCTTCGATGATCAACATGTCAGCAGGATAACACAAGAGGACATTAAGAAAACACATGGTGGATCTTCAG** AATGCATTATATAAGTGGGAATTTGAAGAATCTGAAGAAGATCCAGTGACAAGTATTCCATACCAACTTCAAAGGC -CTTGAAACCAATAGTGGAACTGAAAAGATCTCAAAATCTGGACTTGAAAAGAATTCCTTGATCTATGAACTTTTC1 GCTGGAGGATTCCAGTGCTGGGGAAGACAGTGTTCATGACAGGTTTATAGGTCCGCTTCCAAGAGAAGGTTCTG SAACGTTGTAAGAAGAGGAGGGCGTTCGGTTTTTGCATTTTCCTTATCTGCTGACCTTACAG AAATGAAGGTAGTTGTCACAGTGATCAGATGACCAACGATTTCTCCCAATGATGATGGTGTTGATGAAGGAATCTG ATGGTGCCCGGCGAGGAGCAACTGGTCCCGAAAGAGGCACCACTGGATCATACCAGTGACAAGTCACTTC 4TTTGCTAGTGTGGAAGAAGCATTGCATGCATTTATTCAGCCAGAGATTCTGGATGGCCCAAATCAGTATTTTG1 SGGACTAGTAAACCAAGCAATGACTTGCTATTTGAATAGCCTTTTGCAAACACTTTTTATGACTCCTGAATTTAGG <u> 2TGTTATGGCTCATTCTGGGAGCGCTGCTGGTGGTCATTATTATGCATGTATAAAGTCATTCAGTGATGAGCAGT</u> FGGGTTCTACCAGTGATTATGTCAGCCAAAGCTACTCCTACTCATCTATTTGAATAAATCAGAAACTGGATATGI CGACGCTAATTTTGAGCCAGGAAAGAAGAACTTTCTGCATTTGACAGATAAAGATGGTGAACAACCTCAAATACT

4GGTTGTATTGGAAAGCAGTAGTGTGGACGAATTGCGAGAGAAGCTTAGTGAAATCAGTGGGATTCCTTTGGATG CTGTGGAAGCTATTCTAGAAGAAAGCACTGAAAACTCAAAAGCTTGTCACTGCAGCAACAGCAGGATGGAGATA <u> SAGCATCAGTGGATAATAGAGAACTTGAACAGCATATTCAGACTTCTGATCCAGAAAATTTTCAGTCTGAAGAACG</u> ACAAAACAAGTAATGATGGAAAATAAATTGGAGGTTCATAAGGATAAGACATTAAAGGAAGCAGTAGAAATGGCTT ATATGATGAGAGTGGCAAGAGTAGGGGAGAAATGCAGTACATGTATTTCAAAGCTGAACCTTATGCTGCAGATGA TTAGAGCCCTTTGTTGGAGTTTTGTCCTCTCACTTCAAGGTCTTTCGAGTGTATGCCAGCAATCAAGAGTTTGAG TAGAGCTCAGTATTGACAGGTTTCGTCTAAGGAAAAAAACATGGAAGAATCCTGGCACTGTCTTTTTGGATTATC GTTGGAAGAACAAGAAAAGAGACAACGAGAAATTGAGCGCAATACATGCAAGATAAAATTATTCTGTTTGCATCCT <u> FCTAGAACGGTCATATGAAGGAGAAGAAGATACACCAATGGGGCTTCTACTAGGTGGCGTCAAGTCAACATATAT</u> TACAATGATTTGCGTCTTCTCAGTGTCTCCAGTAAAACCCTGAAAGCTGAAGGATTTTTAGAAGTAACAAGGTGT **ATCAGACTCAGATGTGAATAATGACAGGAGTACAAGTTCAGTGGACAGTGATATTCTTAGCTCCAGTCATAGCAG** TGATACTTTGTGCAATGCAGACAATGCTCAGATCCCTTTGGCTAATGGACTTGACTCTCACAGTATCACAAGTAGT ATAAGATGATGGATTTAGAAGAGGTAATACCCCTGGATTGCTGTCGCCTTGTTAAATATGATGATGAGTTTCATGATTA STTTGATCTGCTGTTGGAGACGAGAAAGCCTGATCAGGTTTTCCAATCTTATAAACCTGGAGAAGTGATGGTGAA TGTTGAAAGCTCCGAGACTTTGGATTACCAGATGGCCTTTGCAGACTCTCATTTATGGAAACTCCTGGATCGGC SAAAGCTGGAGGCGATTCTGGTAATGTGGATGATGACTGTGAAAGAGTCAAAGGACCTGTAGGAAGCCTAAAGT 4GAAGAACGAAAGCAAATGAAGGGAAAAAAAAAACATGGGATACAGCAGAAGAAGACTCTGGAACTGATAGTGA 4GGTTCTGGGGAAGGACATAAATGGTTGATGGTGCATGTTGATAAAAGAATTACTCTGGCAGCTTTCAAACAACA ATATTTATGAAGAAGATATTAATATTTCCAGCAACTGGGAGGTTTTCCTTGAAGTTCTTGATGGGGTAGAGAAGAT SAAGTCCATGTCACAGCTTGCAGTTTTGTCAAGACGGTGGAAGCCTTCAGAGATGAAGTTGGATCCCTTCCAGG SAATCCTAAAGTTTCTACCCTGAATGTCTGGCCTCTTTATATCTGTGATGATGGTGCGGTCATATTTTATAGGGAT ACAGAATTCAAACAACTGATTTCAAAGGCCATCCATTTACCTGCTGAAACAATGAGAATAGTGCTGGAACGCTGC ATATTGAATTTGCTAAGGGTAGAGGAACATTTCCCTGTGATATTTCTGTCCTTGATATTCATCAGGATTTAGACTG **4TGCAAATACAATCAGATTATTTGTTTTGCTACCTGAACAATCCCCAGTATCTTATTCCAAAAGGACAGCATACCA** ATGGGGACAGCAGCAAAAGTACTGAGACAAGTGACTTTGAAAACATCGAATCACCTCTCAATGAGAGGGACTCT **AAAGATCTGACTCAAGACTGA** 

TAAAACGACTGAAGGATGCTTTCAAGAGGACCTGTGGACTCTCATATTACATGGGCCAGCACTGCTTCATCCGGG CAGTTACGTGAATACTACAGAAGAGAAATTTTCAGACAACATTTCTACTGCATCTGAAGCCTCAGAAACTGCTGGC CAGCTGATTCATGGAAGAGACTATGAAATGGTCCCAGAACCTGTGTGGAGGGGCACTTTATCACTGGTATGGAGCA AGTGATACTCCTACTTTCTACCAAACTCTGGCTGGAGTCACACATTTGGAGGAATCAGACATCATTGATCTTGAGA TITITCTAAACAAAGATGCTTTTACTTTCTCTCGATGGCTTCTATCTGGAGGTGTGTATGTTACCCTCACTGATGAT CACCTATTCGTCCATCTCTAAGTGAAGGTTTGTTTAATGCTTTTGATGAAAATCGTGACAATCACATAGATTTTAAG GGACAACCGCACTGATGATATTCCTGAATTACATATGGATCTCTCTGATATTGTAGAAGGCATACTGAATGCACAT ATTATTAGAGGATGGTTAGAACGAGAGAGCAGGTATGGTCTGCAAGCAGGACACAACTGGTTTATCATCTCCATG CTTCTCTTCCTGAGACAGCAGCCTGCCACTCGGACACAGCAGTCTAACATCTGGGGTGAATATGGGAAATGTACCT CACGAATATCTATCTCAAAGGCTGCGCATTAAAGAGGAAGATATGCGCCTGTGGCTATACAACAGTGAGAACTAC CTTACTCTTCTGGATGATGAGGATCATAAATTGGAATATTTGAAAATCCAGGATGAACAACACCTGGTAATTGAAG ATGGGTGCCAAGGAGTCACGGATCGGATTCCTCAGCTACGAGGAGGCGCTGAGGAGAGATTACAGATGTAGAGC GACACCACAAAGATGGGTCATCTTACTCTGGAAGACTATCAGATCTGGAGTGTGAAAAAATGTTGTTGCCAATGAG TTTTGAACCTCCTTTTCCAGGTGTGTCACATAGTTCTGGGGTTAAGACCAGCTACTCCGGAAGAAGAAGAAGACAA CAGTGGTGGCAACAGTGGAAAGAATATGTCAAATACGATGCCAACCCTGTGGTAATTGAGCCATCATCTGTTTTG GTAAAGTCCCAGATACACTCAGGAAGTGTTTCTCAGAGGGTGAAAAGGTAAAACTATGAAAAGTTTAGAAATTGGC AGCGGCTTTCTGTATTCTGCCACACACAGGGGCAGATGTTTGCTTTGCTCGACACATAACACTTCTGACAATAAC <u> AACCAGTGTTTGCTGGGAGCCAATGGGAATATTTTGTTGCACCTTAACCCTCAGAAACCAGGGGCTATTGATAAT</u> CAGCCATTAGTAACTCAAGAACCAGTAAAGGCTACATTAACACTAGAAGGAGGAGGATTAAAAACGAACTCCA TAGTCTTTTTCAAGTGAATCTGGGAACTATGTTATACGGGAAGAAATGGAAAGAATGCTCCACGTGGTGGATG AACGCTATTGGTTATTGAAGGCTCAATCCCGGACTGGACGATTTGATTTAGAGACATTTGGCCCATTGGTTTCAC CCCACAGAAAAGGGAGCCACAGGTCTAAGCAATCTGGGAAACACATGCTTCATGAACTCAAGCATCCAGTGTG AAGTECTTGEGGATGGAGTGCCTCCAAAGGTTGCTGAGGTGATTTACTGTTCTTTTGGTGGAACATCCAAAGGG GAGATATCCTGTGGGTTATCAGCCTGTTGCAGGGGACCCCTGGCTGAAAGACAAAAATTTTGCTTCAAGGTATTT GATGTTGACCGTGATGGAGTTCTCTCCAGGGTTGAACTGAGAGACATGGTGGTTGCACTTTTAGAAGTCTGGAA <u> AATGGAGGAAAATACTCATTTGGAACTGCAGCCCATCCTATGGAGCAGGTCGAAGATAGAATTGGAAGCAGCCT</u> TCTCCGAATGCACCTTTAAAGCGGGTATTAGCCTATACAGGCTGTTTTAGTCGAATGCAGACCATCAAGGAAATT >SGPR480 SEQID 14

STCAGGACAACCAAAAAGTACGACTCTCAGTGAGTGGATTTTTGTGTGCATTTGAAATTCCTGTCCCTGTGTCTCC SAACTTCACAAATGGAATGGTTAATGGTCACATGCCATCTCTTCCTGACAGCCCCTTTACAGGTTACATCATTGCA STGATTATTCACCTTAAGCGATTTCAATTTGTAAATGGTCGGTGGATAAAATCACAGAAAATTGTCAAATTTCCTCG AGAACTTCTGGCTTTTCTCTTGGATGGTCTTCATGAAGATCTTAATCGAGTCCATGAAAAGCCATATGTGGAACTG SATTTGACCCTTTCAATTTTTTGTCTTTGCCACTACCAATGGACAGTTATATGCACTTAGAAATAACAGTGATTAAG STCCACCGAAAAATGATGAGGACAGAACTGTATTTCCTGTCATCTCAGAAGAATCGCCCCAGCCTCTTTGGAATG GCCCCATTAAAGCTTCGGTGGACCATAGCAAAATATGCTCCCAGGTTTAATGGGTTTCAGCAACAGGACTCCCA STGAGTGATCTCTGTGGACTTAATTCAGAACAAATCCTTCTAGCAGAAGTACATGGTTCCAACATAAAGAACTTTC TAGCGAGCCCACTCCCACCTCAGGAAGCTAGTAATCATGCCCAGGATTGTGACGACAGTATGGGCTATCAATAT CCATTCACTCTACGAGTTGTGCAGAAGATGGGAACTCCTGTGCTTGGTGCCCATGGTATAGATTTTGCAGAGGC GTAAAATTGATTGTGGGGAAGACAGAGCTTTCATTGGAAATGCCTATATCGCTGTGGATTGGGATCCCACAGCC CTTCACCTTCGCTATCAAACATCCCAGGAAAGGGTTGTAGATGAGCATGAGAGTGTGGGAGCAGAGTCGGCGAGC ATGTACTACTGTTCCAAGTGTAAGACCCACTGCTTAGCAACAAAGAAGCTGGATCTCTGGAGGCTTCCACCCATC SCATGAAGCATGTGGCAATGGCTACAGCAATGGTCAGCTTGGAAACCACAGTGAAGAAGAAGACGCACTGATGACC rggtatgaaggggcatatggctaaatgctatggtgatttagtgcaggaactttggagtggaactcagaagaatgt GGAAAGTTTTGATCCAAGTGCTTTTTTGGTACCAAGAGCCCGGCTCTCTGCCAGCATAAACCACTCACACCCCA SCATTGATTGTTCCATGTACTGTGCATACCCGGAAGAAGACCTATATGATGCGGTTTGGATTCAAGTATCCGG STTCTTCATCAAGAAAAAGTGGAACCAGCTGTCCCTCCAGCAAAAACAGCAGCCCTAATAGCAGCCCACGGACTI CTGGGGGGGCAGCCAACCAGAGTTGGTCACTCCTCAGGACCATGAGGTAGCTTTGGCCAATGGATTCCTTTATGA CAATTGTTGTGGATTTGTTCCATGGGCAGCTAAGATCTCAAGTAAAATGCAAGACATGTGGGCATATAAGTGTCC SCAAGCCGAGCCCATCAACCTGGACAGCTGTCTCCGTGCTTTCACCAGTGAGGAAGAGCTAGGGGAAAATGAG FGGGGGGGGGCCAAAGGGGGGCTCCGGCTGCCCCAGATTGGCAGCAAAAATAAACTGTCAAGTAAAGAGAA TAGTAACACACAGCCACTGACACAGTATTTATCTCAGGGAGACATCTTTATGAACTCAACAGGACAATCCCA AATGGGGACCTACCCCGACCAATATTCATCCCCCAATGGAATGCCAAACACTGTTGTGCCATGTGGAACTGAGAA GGGGGATGAGCTCTCTGAGCCCAGGATTCTGGCAAGGGAGGTGAAGAAAGTGGATGCGCAGAGTTCGGCTGG CCATTACGTCACTTATGCCAAAAACCCAAACTGCAAGTGGTACTGTTACAATGACAGCAGCTGTAAGGAACTTCA AAGGACAGTGATGGGCGACCAGACTGGGAAGTAGCTGCAGAGGCCTGGGACAACCATCTAAGAAGAAATAGAT

GCCAAAGACTGATGGCAAAAAGATGGCAGACACACAGCAGTATGGATGAAGACTTTGAGTCTGATTACAAAAGTA CCCGGATGAAATTGACACCGACTCTGCCTACATTCTTTTCTATGAGCAGCAGGGGATAGACTATGCACAATTTCT CTGTGTTACAGTAA

>SGPR431 SEQID 15

GCCAGAGTTCGAGTCCTTCTTCAACAAGACCTTCGTGTTGGGCCTCCTTCATCAGGGCTACCACTCTGGACAG AAGTGAGTAACTTGCTGCAGAACATCTGGAAGGCCGAGCCTGCCACACTACTGCCTTCCCTGCAAGAAGTTTTTG TGGTGGAATCGGCGGAGCACTGGCTAGACGAGGCGCAGTGCGAGGCCATGTTTGACCTGACGACCGGCTCAT TCGAACGATAGGCCATTTCCAGTGCGTGTCCACCCAGGAAAGAGAGCTGCGGGAATATGTCTCCCAGGTGACAA SAAGCATCTCCACAGATGCATCATTTGAACCTTCTGTAGCATTGGCAAGCCTTGTGCAGCATATTCCTCTTCA SAGCGACCTTCTGACCGACTTTGTGCAATGCATCCCCAAGGGGAAATTGTCCATCACGTTCTGTCAACAGCTGGT CAGAATGATTGACTGGCTATCCTGGCCATTGGCTCAGCATGTGGATACATGGGTAATTGCACTCCTGAAAGGACT 3GCAGCTGTCCAGAAGTTTACTATTTTGATAGATGTTACTTTGCTGAAAATAGAACTGGTTTTTAATCGACTTTGG1 SAAAACATGTCTGTCCAAGATCCAGCATCATCACCCAGTATACAAGATGGTGGTCTAATGCAAGCCTCTGTACCC CCTGGAGGGCCAGGACCCTTTCCAGCGGCAGGTGGGGCACCAGGTGCTGGAGGCCTACGCACGATACCACCG GAAGGATGTAGCCATCCTGGACTACATTCACAACGGCCTGAAGCTGATTATGAGCTGTCCGTCGGTGCTGGATC <u> CCATTTGATTGTTCCTCATGTGGTTAATTTGGTTCATTCTTTCAAAAATGATGGTCTGCCTTCAAGTACAGCCTTCT</u> TAGTACAATTAACAGAATTGATACACTGTATGATGTATCATTATTCTGGATTTCCAGATCTCTATGAACCTATTCTG STGAATACCTCAGATTTCTCCTTGACAGGCTCCATGAAGAAGAAAAGATCTTGAAAGTTCAGGCCTCACACAAGC CTCGTACAAGTGACGGTGAGAAGACTTTAATAGAAAAAATGTTTGGAGGAAAACTACGAACTCACATACGTTGTTT TTCCTCTTGTGAGACCTGGTGCTCTTTGCAGTTCTTTCTCACATGCTGCTTAGCTTTCAGCATTCTCCAGAGGCGTT CTTCTGAAATTCTGGAATGCAGTGAAACTTCTTTACAGGAAGTAGCTAGTAAAGCAGCAGTACTAACAGAGACCC ATGGACAAGATCCTGGAGGGCCTTGTGAGTTCCTCGCATCCCCTGCCCCTCAAGCGGGGTGATTGTGCGGAAGG TCTTTAGCCTCCTGCAGGTAGAGGTGTTACGGATGGTGTGTGAGAGGCCGGAGCCGCAGCTCTGTGCCCGACT GATGATTACAGTTCTCATCAGGAGCCTTACTACGGATCCAAATGTAAAAGATGCAAGTATGACCCAAGCCCTTTG ITAAATCTAAATGGGTGCAATTCATTAATGAAAAATTACAGCATCTTTTTGCCTTTCTGGCCCATACACAGAGGG <u> AATCCAATTCTTTGGCGTCTTGCTTGTCTAGACTTTCTGGAAAATCTGAAACTGGGAAAACTGGTCTTATTAACCT</u> <u> AGGAAATACATGTTATATGAACAGTGTTATACAAGCCTTGTTTATGGCCACAGATTTCAGGAGACAAGTATTATCT</u> **AAGCATACGCACCTCGGATATTCTTTGAGGCTTCCAGACCTCCATGGTTTACTCCCAGATCACAGCAAGACTGT** 

FIG. 1R

CAGAAAGTTGGTCTGTAGATGTTGACTTCACTGATCTTAGTGAGAACCTTGCTAAAAAATTAAAGCCTTCAGGGAC <u> AGTGGGCATTACTATTCTTATGCCAGAAATATCACAAGTACAGACTCTTCATATCAGATGTACCACCAGTCTGAGG</u> CTCTGGCATTAGCATCCTCCCAGAGTCATTTACTAGGGAGAGATAGTCCCAGTGCAGTTTTTGAACAGGATTTGG SCAGAGATTCTTACTGGTGATAACCAATATTATTGTGAAAACTGTGCCTCTCTGCAAAATGCTGAGAAAACTATGC I GATGAAGCTTCCTGCACAAAATTGGTGCCCTATCTATTAAGTTCCGTTGTGGTTCACTCTGGTATATCCTCTGAA CATGTTCATTTCGGCCCAATGGATTTGATGACAACGACCCACCAGGAAGCTGTGGACCAACTGGTGGAGGGGGGT **AAATCACGGAGGAACCTGAATACCTTATTCTTACTCTCCTGAGATTTTCATATGATCAGAAGTATCATGTGAGAAG AAAATTACGAGCAGGTTTCCAAAGGACACAGCTTATGTGCTTTTGTATAAAAAACAGCATAGTACTAATGGTTTAA** CAAAAGACAATAAACTATATTACAGGAACAAGAGTTGAATGCTCGAGCCCGGGCCCTCCAAGCTGCATCTGCTT 7AATAAAGATGTACCTCAGAAACCAGGAGGTGAAACCACACCTTCAGTAACTGACTTACTAAATTATTTTTGGC1 <u> CCATAGGCAGTCCTCCTAATGAGTTTTACTGTTCTGAAAACACTTCTGTCCCTAACGAATCTAACAAGATTCTTG</u> **3GAGGAGGATTTAATACAGTTGGCAGACTCGTATTTTGA** 

STACTTGTTTGCCTTGTTGCAAAACAGTAATAGGCGATACATTGATCCATCAGGATTTGTTAAAGCCTTGGGCCTG &AAAGAATCCAGATGTGCGCAATATTGTTCAACAGCAGTTCTGTGGAGAATATGCCTATGTAACTGTTTGCAACCA GTGTGGCAGAGAGTCTAAGCTTTTGTCAAAATTTTATGAGCTGGAGTTAAATATCCAAGGCCACAAACAGTTAACA SATTGTATCTCGGAATTTTTGAAGGAAGAAAATTAGAAGGAGACAATCGCTATTTTGCGAGAACTGTCAAAGCA SGCCACTACATCGCCCACGTGAAAGATCCACAGTCTGGTGAATGGTATAAGTTTAATGATGAAGACATAGAAAAG 
 FAGTGACTACATGCTGGGAGACGGCATCCAAGAAGAAAAAGATTATGAGCCTCAAACAATTTGTGAGCATCTCCA
'GACAGGCAAACTGGACATAAGAAAAAGCTGAATACCTACATTGGCTTCTCAGAAATTTTGGATATGGAGCCTTAT SACACTGGACAACAGCAGGATGCTCAAGAATITTCAAAGCTCTTTATGTCTCTATTGGAAGATACTTTGTCAAAC GCAGGAGCATATCGAGACCGCTTACCGCATCTGGCTGGAGCCCTGCATTCGCGGCGTGTGCAGACGAACTGC 4CATCGATGATCCCAACTGTGAGAGGAGAAAAAAAACTCATTTGTGGGCCTGACTAACCTTGGAGCCACTTGT >SGPR429 SEQID 16

ATGGAGGGAAGAAATTACAACTAGGGATTGAGGAAGATCTAGAACCTTCTAAGTCTCAGACACGTAAACCCAAG TGTGGCAAAGGAACTCATTGCTCTCGAAATGCATATATGTTGGTTTATAGACTGCAAACTCAAGAAAAGCCCAACA CTACTGTTCAAGTTCCAGCCTTTCTTCAAGAGCTGGTAGATCGGGATAATTCCAAATTTGAGGAGTGGTGTATTGA TAAAAGAATTGAAAATTCAGATCATGCATGCATTTTCAGTTGCTCCTTTTGACCAGAATTTGTCAATTGATGGAAAG ATTITAAGTGATGACTGTGCCACCCTAGGCACCCTTGGCGTCATTCCTGAATCTGTCATTTTATTGAAGGCTGATG GGTTACCTGCTGGAGCTGAGCCCTATGAGTTTGTCTCTCTGGAATGGCTGCAAAAGTGGTTGGATGAATCAACAC CAGAAACACAGTATATTTCTGAGCCCAAACTCTGTCCAGAATGCAGAGGAAGGCTTATTGTGTCAGCAGCAGAGGG TCCAAAGGCTCCTGAGTTTCCAAGTTACAAAGAGTGCTGTTCACAGGGCAAGATTTTAGAAAGAGAGGGGAAGA GAAATTTGTTAGAAAGCCTACAAGATGCAGCCCTGTGTCATCAGTTGGGAACAGTGCTCTTTTGTGTCCCCACGG CTACCAAACCTATTGATAATCACGCTTGCCTGTGTTCCCATGACAAGCTTCACCCGGATAAAATATCAATTATGAA TAAATAAAGATGAATCAAAGGAAGAAAGAAAGAAGAGGGGGAGTTAAATTTAATGAAGATATTCTGTGTCCACA AAATGAAGCCTTACATAAGATGATTGCAAACGAGCAAAAGACTTCTCCCCAAATTTGTTCCAGGATAAAAACAGA AATGGCTGAGATGCGTAAGCAAAGTGTGGATAAAGGAAAAGCAAAACACGAAGAGGGTTAAGGAGCTGTACCAAA GCTAGCTCTTGAACAGCTGGATGAGCAAGATGGTGATGCAGAACAAAGCAACGGAAAGATGAACGGTAGCACCT ACCTGCGTGAATACACTCAAGCCACCATCTATGTCCATAAAGTTGTGGATAATAAAAAGGTGATGAAGGATTCGG AACCAATTGCAGATTATGCTGCAATGGATGATGTCATGCAAGTTTGTATGCCAGAAGAAGGGTTTAAAGGTACTG CTCCGGAACTGAATGTGAGTAGTTCTGAAACAGAGGAGGACAAGGAAGAAGCTAAACCAGATGGAGAAAAAGAT GTGTAAGGAATGTGTAGTAGAACGTTGTCGCATATTGCGTCTGAAGAACCAACTAAATGAAGATTATAAAACTGT CAAATGATACAAAAGCTCTTTGTTGTGGATCATGTAATTAAAATCACGAGAATTGAAGTGGGGAGATGTAAACCCTT SAGGATATCTGAATATGCAGCTGACATTTTCTATAGTAGATATGGAGGAGGTCCAAGACTAACTGTGAAAGCCCT CCGTGTCTCAGTAACTGGCCAGAGGATACGGATGTCCTCTACATCGTGTCTCAGTTCTTTGTAGAAGAGTGGCG <u> AAGTTÄTTCGCCGAAGTATGCGACATAGAAAAGTTCGTGGTGAGAAAGCACTTCTCGTTTCTGCTAATCAGACGT</u> TGGTGAGTTATGCATATCTGAAAATGAAAGAAGGCTTGTTTCTAAAGAGGCTTGGAGCAAACTGCAGCAGTACTI

FIG. 1T

>SGPR503 SEQID 17

ATECTGAGCTCCCGGGCCGAGGCGGCGATGACCGCGGCCGACAGGGCCATCCAGCGCTTCCTGCGGACCGGG ATATGTTATTTGGGGTCCCATTACAGAAAGAAGGAAGCGTAGAAAGGGGCTTGTGCCTGGCCTTGTTAATTTAGG GCGGCCGTCAGATATAAAGTCATGAAGAACTGGGGAGTTATAGGTGGAATTGCTGCTGCTCTTGCAGCAGGAAT

GTACAGATGGCAGATCTCATCTTGAAGAACAGGATGCTCACGAATTATTCCATGTCATTACCTCGTCATTGGAA AGCCTTTCACTAAGTATTCCAGCCGCCACATGGGGTCACCCATTGACCCTGGACCACTGCCTTCACCACTTCATC ICATGGAAGACTCACTAGTAATATGGTCTGCAAACACTGTGAACACCAGAGTCCTGTTCGATTTGATACCTTTGAT TCATCAGAATCAGTGCGGGATGTTGTGTGTGACAACTGTACAAAGATTGAAGCCAAGGGAACGTTGAACGGGGA AAAGGTGGAACACCAGAGGAGCACTTTTGTTAAACAGTTAAAACTAGGGAAGCTCCCTCAGTGTCTCTGCATCCA GTGGCTGTGGGTCTCCGATGACACTGTCCGCAAGGCCAGCCTGCAGGAGGTCCTGTCCTCCAGCGCCTACCTG CCTACAGCGGCTGAGCTGGTCCAGCCACGGCACGCCTCTGAAGCGGCATGAGCACGTGCAGTTCAATGAGTTC CATGCACTCTGGACACTTTGTCACTTACCGACGGTCCCCACCTTCTGCCAGGAACCCTCTCTCAACTAGCAATCA <u> GAACACCTGCTTCATGAACTCCCTGCTACAAGGCCTGTCTGCCTGTCCTGCTTTCATCAGGTGGCTGGAAGAGT</u> SAAAGCCTTGTCCTGCCAAGAAGTTACTGATGATGAGGTCTTAGATGCAAGCTGCTTGTTGGATGTCTTAAGAAT CCAGGGCCTACACTGGAGCTGCAGGATGGGCCGGGAGCCCCCACACCAGGTTCTGAATCAGCCAGGGGCCCCC AAAACACAGATTTTTATGAATGGCGCCTGCTCCCCATCTTTATTGCCAACGCTGTCAGCGCCGATGCCCTTCCCT CACCTCCCAGTACTCCAGGGATCAGAAGGAGCCCCCCTCACACCAGTATTTATCCTTAACACTCTTGCACCTTCT TCCCAAACAAATTACCTGCCGCACAAGAGGGTCACCTCACCCCACATCCAATCACTGGAAGTCTCAACATCCTT CTCCCAGTTGTTCCCGACTACAGCTCCTCCACATACCTCTTCCGGCTGATGGCAGTTGTCGTCCACCATGGAGA CTGTTCTACGAGCGCGTCCTTTCCAGGATGCAGCACCAGAGCCAGGAGTGCAAGTCTGAAGAATGA

CCAGCCGGGACACTGTGATGGCGACGGTGAGGGGGGGTTCGCCTGCGCCCCGGGCCCAGTTCCAGCGGCCCC CGAGCAGCTGGCGCGCTGGTGCGCGCGCTCTGGACTCGCGAATACACGCCCCAACTTTCCGCGGGAGTTCAAG AATGCAGTTTCCAAGTACGGCTCTCAGTTCCAAGGCAATTCCCAGCACGACGCCCTGGAATTCCTGCTCTGGTTG CGGGAGCCCCGGGGAGGAACGCCCGGCCCGGACCCCAGCCCCAGCTCCAGCTCCCGCCGGCGGGGGGCGC GGCCGCCGGGCGCTCAGGGCTTGAAGAACCACGGCAACACCTGTTTCATGAACGCGGTGGTGCAGTGTCTCAG GCGCCGCCTGTTCAGCCGCTTCCTGCTGGCGCTGGGCAGCCGCTCACGCCCCGGGGGACTCACCGCCCGGGCC CTGGATCGTGTACATGAGGACCTGGAGGGTTCATCCCGAGGGCCGGTGTCGGAGAAGCTTCCGCCTGAAGCCA <u> AAGCACAATATAGATCTTCCTTGACTTGTCCCCACTGCCTGAAACAGAGCAACACCTTTGATCCTTTCCTGTGTGT</u> CTAAAACCTCTGAGAACTGCCTGTCACCATCAGCTCCAGCTTCCTCTAGGTCAAAGCTTTGTGCAAAGCCACTTTC STCCCTACCTATCCCCTTGCGCCAGACGAGGTTCTTGAGTGTCACCTTGGTCTTCCCCTCTAAGAGCCAGCGGT >SGPR427\_SEQID\_18

 
 FCCTGCGGGTTGGCCTGGCCGTGCCGATCCTCAGCACAGTGGCAGCCCTGAGGAAGATGGTTGCAGAGGAGG
CTCCACAGTGAGGAGGAGGTGCTAATCCTCTTCTGTAACTTGGTGGGGTCAGGGCAGCAGGCTAGCAGGTTTGG GGGAGCCTCCAGGAGGAGCGAGCGCAGGATGCCGACAGTGTGTGGCAGCAGCAGCAGCAGGCGCATCAGCAGCAC 
 AAGAAGGAGAACAGGAGGAATGAGAGGGCAGAGGTCTCTCCACAGGTGCCCCCCCGTCTCCCTGGTGAGTGGCG
CAGGAGGCCAGGAGGCCCTCCACATGTCAAGCTGGCGGTGGAGTGGGATAGCTCTGTCAAGGAGGGCCCTGTTC TTACGGCTCGGGAGCCACGCTGGCAGCACAAGGGGAAGCCTGCTGTCCTGGAGCTCTGCCCCCTGCCCTCCC <u> ICTCAGCTCATCCACTGGGTCTGTCGGCCTCCCCACGCCTGGCAGCCCGTGAGGGCCAGCGATTCTCCCTCTCT</u> GCCACCCTTCCTGATAAGGGAAGACAGAGCTGTTTCCTGGGCCCAGCTCCAGCAGTCTATCCTCAGCAAGGTCC AGCTGTACCTTGGATGATGTTTTCAGTTCTACACCAAGGAGGAGCAGCTGGCCCAGGATGACGCCTGGAAGTG **ACCTCAAAAGGTTCTGCCAGGTGGGCGAGAGAAGAACAAGCTCTCCACGCTGGTGAAGTTTCCGCTCTCTGGA** CTCAACATGGCTCCCCATGTGGCCCCAGAGAAGCACCAGCCCTGAGGCAGGACTGGGCCCCTGGCCTTCCTGGA AGCAGCCGGACTGCCTGCCCACCAGTTACCCGCTGGACTTCCTGTACGACCTGTATGCCGTCTGCAACCACCAT GGCAACCTGCAAGGTGGGCATTACACAGCCTACTGCCGGAACTCTCTGGATGGCCAGTGGTACAGTTATGATGA CAGCACGGTGGAACCGCTTCGAGAAGATGAGGTCAACACCAGAGGGGCTTATATCCTGTTCTATCAGAAGCGGA CATTCAAGACCATGCCTCTGCGGTGGTCCTTTGGATCCAAGGAGAAACCACCAGGTGCCTCCGTCGAGTTGGTG STGAGGATGAGAAGTCAGCATCGCCGAGGTCCAACGTCGCCCTTCCTGCTAACAGCGAAGATGGTGGGGGGG TCCTCACTGCCAAGTCCTGCAGGGGATGGTGAAGCTGAGTTTGTGGACGCTGCCTGACATCCTCATCATC ACAGCATCCCTCCCTGGTCAGCCAGCAGCTCCATGAGAGGCTCTACCAGCTCCTCCTGCTGTCTGATCACTGGCTC AAGGAGAGCTTCCAGATGGGAAGCAAAAGCAGCCCACCCTCCCCCTATATGGGATTCTCTGGAAACAGCAAAGA GCCTCTCACTGTGATGCCTTCAGTGGAGCATGAGAACCAGCTCGACCGGAGGGCCAGAAGGCCATGAACTGG GTGGCCTGCAGCTATTTGTCTCCGAAGGACAGTCGGCCCCTCTGTCACTGGGCAGTTGACAGGGTTTTGCATCT SAGTACTTGGAATCCAGACGAAGACCTCGGTCCACGAGCCAGTCCATTGTGTCGCTGTTGACGGGCACTGCGG CATTGAAAGAGGTCCAGCCGGGGTGCCCTGTCCCTCGGCTCAACCCAACCACTGTCTGGCCCCTGGAAACTCA CAGTCGCCGAGGCACCTCTGAGCTAGACAGACCCCTGCAGGGGGACACTCACCCTTCTGAGGTCCGTGTTTCGG SGCTGAGCCCTGCCATGGACGGGCAGGCTCCAGGCTCACCTCCTGCCCTCAGGATCCCAGAGGGCCTGGCCA SATGGTCCAAACACAGGAAGGAAACTCAAGGAAAATGCAGGGCAGGACATCAAGCTTCCCAGAAAGTTTGACCT GCCATCTTATGAAGAGTGAGGCCCCTGTACAGAACCTGGGGTCTCTGTTCTCCATCCGTGTTGTGGGACTCTCT SGGGCCTGGGCAGCCGGCTCGAGAGGGATGTCTGGTCAGCCCCCAGCTCTCCGGCCTCCCCCCTCGTAAAGCCA GCAGGGCCCCGAGAGGCAGTGCACTGGGCATGTCACAAAGGACTGTTCCAGGGGAGCAGGCTTCTTATGGCAC 

>SGPR092 SEQID 19

ATAATCTGTTCCTGGTTGCAACTCATGAGATTGGGCCATTCTTTGGGCCTGCAGCACTCTGGGAATCAGAGCTCCA TAATGTACCCCACTTACTGGTATCACGACCCTAGAACCTTCCAGCTCAGTGCCGATGATATCCAAAGGATCCAGC AATAAGCACACTCTAACTTACAGGATTATCAATTACCCACATGATATGAAGCCATCCGCAGTGAAAGACAGTATAT ATGCAGCTCGTCATCTTAAGAGTTACTATCTTCTTGCCCTGGTGTTTCGCCGTTCCAGTGCCCCCTGCTGCAGAC ATAATGCAGTTTCCATCTGGAGCAATGTGACCCCTTTGATATTCCAGCAAGTGCAGAATGGAGATGCAGACATCA AGGTTTCTTTCTGGCAGTGGCCCCATGAAGATGGTTGGCCCTTTGATGGGCCAGGTGGTATCTTAGGCCATGCC TTTTACCAAATTCTGGAAATCCTGGAGTTGTCCATTTTGACAAGAATGAACACTGGTCAGCTTCAGACACTGGAT TCTGCTACACCAGCCCCACTGTGGGGTGCCTGATGGGTCCGACACCTCCATCTCGCCAGGAAGATGCAAGTGG CATAAAGGATGGGACTTTGTTGAGGGCTATTTCCATCAATTTTTCCTGACCAAGAAGGAGTCGCCACTCCTTACC ATTTGTATGGAGAAAAATGTTCATCTGACATACCTTAA

>SGPR359\_SEQID\_20

AAATAAAGAAGGACACCAGATTGGTGAGATGGTTGCAAGAGGAAGCAATTCCATGATAAGGAAGATTAAGGAGCT GTACAAGAATCCCTATGGATTCCACCTCCCCAAAGATGATGTGAAAGGGATCCAGGCATTATACGGACCTCGGAA FIGOTICAT GAATT TIGG COAT GOOC TIGG COOT GOOC CATTIC CACAGA COCATICA GOOC TIGATION TO THE ACTIVITY AND A COLOR OF THE ACTIVITY <u> ATGAAGGTGCTCCCTGCATCTGGCTTGCTGTTTCCTCATCATGGCTTTGAAGTTTTCCACTGCAGCCCCTCC</u> GATCACGGGGATTCCTATCCATTCGATGGGCCTCGGGGGACTCTAGCCCATGCATTTGCTCCTGGAGAGGCCT GGGAGGAGATACACATTTCGACAATCCTGAGAAGTGGACTATGGGAACGAATGGTTTTAATTTGTTTACCGTTGC SCTCATCCTTTGACGCTGTGACAATGCTGGGGAAGGAGCTCCTGCTCTTCAAGGACCGGATTTTCTGGAGACGG CTAGTTGCAGCCTCCCCCAGGACCTGGAGGAACAACTACCGCCTCGCACAGGCGTATCTTGACAAATATTACAC GAATATCTAAATACACACCTTCCATGAGTTCTGTCGAGGTGGACAAAGCAGTGGAGATGGCCTTGCAGGCCTGG AGTAGCGCCGTCCCTCTGAGCTTTGTCAGAATAAACTCAGGAGAGGGGGATATTATGATATCTTTTGAAAATGGA CAGGTTCACTTGCGGACAGGAATTCGGCCCAGCACTATTACCAGCTCCTTCCCCCAGCTCATGTGGGA ACAAGCGTTCTTTGGCCTCCAAGTCACCGGGAAGTTAGACCAGACCACAATGAACGTGATCAAGAAGCCTCGCT TGCAGCTTACGAAGTGGCTGAGAGGGGCCACTGCTTACTTCAAAGGTCCCCACTACTGGATAACAAGAGGAT rccaaatgcaaggtcctcctcgactatttatgactttggatttccaaggcacgtgcagcaaatagatgctgctg ICTACCTCAGGGAGCCACAGAAGACCCTTTTCTTTGTGGGAGATGAATACTACAGCTACGACGAAAAGGAAAAGG **AAAATGGAAAAAGACTATCCAAAGAATACTGAAGAAGAATTTTCAGGAGTAAATGGCCAAATCGATGCTGCTGTA** STGTGGTGAAATCTAGTTCCTGGATTGGTTGCTAA

>SGPR104\_1\_SEQID\_21

STGAACTTCGGTGGCATCGGTGTGGTCATGGGCCATGAGTTGACGCATGCCTTTGATGACCAAGGGGCGCGAGTA <u> ATGAACGTCGCGCTGCAGGAGCTGGGAGCTGGCAGCAACATGGTGGAGTACAAACGGGCCACGCTTCGGGATG</u> <u> S<u>G</u>CTTGAGTTCCTGTCTTTCTTGCTGTCACCATTGGAGTTGAGTGACTCTGAGCCTGTGGTGGTGTATGGGATGG</u> <u> GGGGTCCCTCTTCGTGAAGGCCACGTTTGACCGGCAAAGCAAAGAAATTGCAGAGGGGATGATCAGCGAAATCC</u> <u> AAGACGCACCCGAGACCCCCGTAGAGGGCGGGGCCTCCCCGGGACGCCATGGAGGTGGGATTCCAGAAGGGGA</u> <u> CACGAGGGAGCAGATGCAGCAGGTGCTGGAGTTGGAGATACAGCTGGCCAACATCACAGTGCCCCAGGACCAG</u> ATTATTTGCAGCAGGTGTCAGAGCTCATCAACCGCACGGAACCAAGCATCCTGAACAATTACCTGATCTGGAACC CAACTAAGAATGAGATCGTCTTCCCCGCTGGCATCCTGCAGGCCCCCTTCTATGCCCGCAACCACCCCAAGGCC TGGTGCAAAAGACAACCTCAAGCCTGGACCGACGCTTTGAGTCTGCACAAGAGAAGCTGCTGGAGAGCCTCTAT <u> ACAGCAATGTTATCCAGGTGGACCAGTCTGGGCTCTTTCTGCCCTCTCGGGATTACTACTTAAACAGAACTGCCA</u> CGGCGCGACGAGGAGAAGATCTACCACAAGATGAGCATTTCGGAGCTGCAGGCTCTGGCGCCCTCCATGGACT GTTCTCCTGTGGGGGCTGGATTCGGAGGGAACCCCCTGCCCGATGGGCGTTCTCGCTGGAACACCTTCAACAGC SCAGAAGACACAGGGCTTCTACCTATCTTGCCTACAGGTGGAGCGCATTGAGGAGCTGGGAGCCCAGCCACTG SAAGGCAGTAGCAGGGACCTACAGGGCCACCCCATTCTTCACCGTCTACATCAGTGCTGACTCTAAGAGTTCCA AGCAGATGCCATCTATGATATGATTGGTTTCCCAGACTTTATCCTGGAGCCCAAAGAGCTTGGATGATGTTTATGA SACCAGCTCCGCAAGCCTCCCAGCCGAGACCAGTGGAGCATGACCCCCCAGACAGTGAATGCCTACTACCTTC 4GAGACCTCATTGAGAAGATTGGTGGTTGGAACATTACGGGGCCCTGGGACCAGGACAACTTTATGGAGGTGTT CCTGCATTCGAGTGGCTGGAAAAATCCTGGAGTCCCTGGACCGAGGGGTGAGCCCCTGTGAGGACTTTTACCA <u> GECACTAAGAAGTCCTGTGTGCCGAGGTGGCAGACCTGCATCTCCAACACGGATGACGCCTTGGCTTTGCTT</u> CGGGTACGAAATTTCTGAAGATTCTTCTTCCAAAACATGTTGAATTTGTACAACTTCTCTGCCAAGGTTATGGCT

FIG. 1X

ACTCTCTCCAACTCCCGTGACTTCCTGCGGCACTTCGGCTGCCCTGTCGGCTCCCCCATGAACCCAGGGCAGCT TGACAAAGAAGGGAACCTGCGGCCCTGGTGGCAGAATGAGTCCCTGGCAGCCTTCCGGAACCACACGGCCTGC ATGGAGGAACAGTACAATCAATACCAGGTCAATGGGGAGAGGCTCAACGGCCGCCAGACGCTGGGGGAGAACA TTGCTGACAACGGGGGCTGAAGGCTGCCTACAATGCTTACAAAGCATGGCTGAGAAAGCATGGGGAGGAGCA GCAACTGCCAGCCGTGGGGCTCACCAACCAGCTCTTCTTCGTGGGATTTGCCCAGGTGTGGTGTGCTCGGTC GTGTGAGGTGTGGTAG

# >SGPR303 SEQID 22

CTGGGACCTTGTTACTTATAGGGAGACTGCATTGCTTATTGATCCAAAAAATTCCTGTTCTTCATCCGCCAGTGG GGATTTAACTATCAGAATGAAGATGAAAAAGTCACCTTGTCTTTCCCTAGTACTCTGCAAACAGGTACGGGAACCT 36ACTCAGTTTGTTTCTGCTGATTACACCCGTGCCCAGGAGCTTGACGCCTTAGATAACAGCCATCCTATTGAAG AGAGGTGCGCTATGCTGCTGTAACACAGTTTGAGGCTACTGATGCCCGAAGGGCTTTTCCTTGCTGGGATGAGC ACTCCTGTTGGCAAAGCAGAACAAGGAAAATTTGCATTAGAGGTTGCTGCTAAAACCTTGCCTTTTTATAACGACT TGTGATGAATTGTGCTGATATTGATATTACAGCTTCATATGCACCAGAAGGAGATGAAGAAATACATGCTACA GTGCTATCAAAGCAACTTTTGATATCTCATTGGTTGTTCCTAAAGACAGAGTAGCTTTATCAAACATGAATGTAATT <u> ACTTCAATGTTCCTTATCCTCTACCTAAAATTGATCTCATTGCTATTGCAGACTTTGCAGCTGGTGCCATGGAGAA</u> GACCGGAAACCATACCCTGATGATGAAATTTAGTGGAAGTGAAGTTTGCCCGCACACCTGTTACATCTACATAT GTTGCTCTGGTTGTGGGACATGAACTTGCCCATCAATGGTTTGGAAATCTTGTTACTATGGAATGGTGGACTCAT 1CAGTGTGGGCCATCCATCTGAGGTTGATGAGATATTTGATGCTATATCATATAGCAAAGGTGCATCTGTCATCC ATGCCGGAGAAGAGGCCCTTCGAGCGGCTGCCTGCCGATGTCTCCCCCATCAACTGCAGCCTTTGCCTCAAGC <u> CTTTGGTTAAATGAAGGTTTTGCATCCTGGATTGAATATCTGTGTGTAGACCACTGCTTCCCAGAGTATGATATT</u> 

FIG.

**1Y** 

# >SGPR402 1 SEQID 23

GTCGAGCGCCTGGCACGCAAATTCGGCTTCGTCAACCTGGGGCCGATCTTCCCTGACGGGCAGTACTTTCACCT GGGTGGGCCCCGGTCCGAGCCCCCATCTATGTCAGCAGCTGGGCCGTCCAGGTGTCCCAGGGTAACCGGGAG SCGGCACCGGGGCGTGGTCCAGCAGTCCCTGACCCCGCACTGGGGCCCACCACCTGCACCTGAAGAAAACCC ATGCGGCCCGGATTGCGCTGTGGCCTGCGCCTGGTCTTGGCCCTGGCCCTTGTCCGCCCCGGGCTGTG

AACAGCATCCACACGCTTTCCGTGGGCAGCACCACCCAGGAGGGCCGCGTGCCCTGGTACAGCGAAGCCTGC GCTACAACAACTGGGTCTTCATGTCCACCCACTTCTGGGATGAGAACCCACAGGGCGTGTGGACCCTGGGCCTA GGGCACGCTCTTCATCTGGGCCTCGGGCAACGGCGGCCTGCACTACGACAACTGCAACTGCGACGGCTACACC GCAGGCCGAGGACTGGAGGACCAACGGCGTGGGGCGCCCAAGTGAGCCATCACTACGGATACGGGCTGCTGGA CCACACAAGGCTGGTGACCGCTGGGCCTGGGCACACGGCGCGCCCCGCGCTGAGGGTCTGCTCCAGCTGCCA TGCCTCCTGCTACACCTGCCGCGGCGGCTCCCCGAGGGACTGCACCTCCTGTCCCCCATCCTCCACGCTGGAC CACGGTGGACGGCCCCGGCATCCTCACCCGCGAGGCCTTCCGGCGTGGTGTGACCAAGGGCCGCGGGGGCT TGGGGGTCGCTTTCAACGCCCGAATCGGAGGCGTACGGATGCTGGACGGTACCATCACCGATGTCATCGAGGC CCTCCACCCTCACCACCTACAGCAGCGGCGTGGCCACCGACCCCCAGATCGTCACCACGGACCTGCATCA CCAGAGCCGCCCCACCCCATCCTGCCGCTGATCTACATCAGGGAAAACGTATCGGCCTGCGCCGGCCTCCAC GCAAAGAGACCGGCACGGGACCCGCTGTGCTGGGGAGGTGGCCGCGATGGCCAACAATGGCTTCTGTGGTG AACTCCATCCGCTCGCTGGAGCACGTGCAGGCGCAGCTGACGCTGTCCTACAGCCGGCGCGGAGACCTGGAG GAGAACAAGGGCTACTATTTCAACACGGGGACGTTGTACCGCTACACGCTGCTGCTGTATGGGACGGCCGAGG ACATGACAGCGCGGCCTACAGGCCCCCAGGTGACCAGCAGCGCGTGTGTGCAGCGGGGGACACAGAGGGGGCTGT CAGCAGCAGGGCTCCTGCATGGGACCCACCACCCCGGACAGCCGCCCCCGGCTTAGAGCTGCCGCCTGTCCC CTGGTTCTCCAAGCAGTGGTACATGAACAGCGAGGCCCAACCAGACCTGAGCATCCTGCAGGCCTGGAGTCAG CCAGTCGCTGAGCCTGCAGCCGCAGCACATCCACATTTACAGCGCCCAGCTGGGGTCCCGAGGACGACGGCCG ATCTCGCTCACCAGCCCCATGGGCACGCGCTCCACACTCGTGGCCATACGACCCTTGGACGTCAGCACTGAAG CAAGGTGCAGTGGTTCCAGCAGCAGAGGCTGCAGCGGGGGGGTGAAACGCTCTGTCGTGGTGCCCACGGACCC GGGCTGTCAGGCCAGGGCATCGTGGTCTTGTGCTGGACGATGGCATCGAGAAGGACCACCCGGACCTCTGG GCCAACTACGACCCCCTGGCCAGCTATGACTTCAATGACTACGACCCGGACCCCCAGCCCCGGTACACCCCCA CACCACCECTECCCAGCCTCGGCCATGGTGCTGAGCCTCCTGGCCGTGACCCTCGGAGGCCCCGTCTTGC GGCATGTCCATGGACCTCCCACTATACGCCTGGCTCTCCCGTGCCAGGGCCACCCCCACCAAACCCCAGGTCT GGCTGCCAGCTGGAACCTGA

1Z

FIG.

GGCCCGGGCAGGCAGGGTGCGCAGGGAGGCGTAGCACTGCTCTTCCCCTCCGCGCTCCCCTCAGGGCC >SGPR434 SEQID 24

LOGGOOTS CLOSEOT

TTGGGTGACTGGCTGGAGCCAGGTTAAGCAGCGCTTTTCAGGCTCCACAGCCAACTCCATGCTGACCCCAGAGC STCACCTTCAGTGAGTACGTGCAGCCCATCTGCCTCCCGGAGCCCAATTTCAACCTGAAGGTTGGGACGCAGTG GTGTGGAGGGGCCCTCATTGACCCCAGCTGGGTGGTGACTGCGGCCCACTGCAGGCCAAGGCCAAAGAGTA **ATCATTATGCACCCCAAGTACTGGGGCCGGGCCTTCATCATGGGTGACGTTGCCCTTGTCCACCTTCAAACACC** <u> AGCCTTCTCAGGTCCCTGTGCCTCTGCCTGCTCCTGTTCCTGTGCTGGCCGCTGCAGCCCCAGATGGGCTCCT</u> CCCAGTCCAGGGGGAACGCAGTGCGCCCCGCTTCGGTGTTACTTCCCTCAGCCTGTGGCCAGCGGGACTTCAAG TGGTGGTCGGAGGATTTGGAAATGACCCGCCATTGGCCCTGGGAGGTGAGCCTCCGGATGGAAAATGAGCACG ATCTGCGGAGGCTCCCTTATCGCCAGACACTGGGTTATAAAGAGGACACAACCAATCCAGTTTGTGGTGAGCCC CTCAGTGGTGCTTGGCACCTCCAAGCTGCAGCCCATGAACTTCAGCAGGGCCCTCTGGGTCCCTGTGAGGGAC FGCAGGAGGCTGAGGTGTTTATCATGGACAACAAGAGGTGTGACCGGCATTACAAGAAGTCCTTCTTCCCCCCA GTTGTCCCCCTTGTCCTGGGGGGACATGATCTGTGCCACCAATTATGGGGAAAACTTGTGCTATGGGGATTCTGG AAGGCACAGAATCCAGGTGTGTACACCCGCATCACCAAATACACCAAATGGATCAAGAAGCAAATGAGCAATGG

>SGPR446 1 SEQID 25

STGACGGCAGGCCACTGCATTTCCAGCCGTTTCCATTACAGTGTCAAGATGGGAGATCGGAGTGTCTATAATGAA TTGCATCAGAAATTCTTCAGGATGTGGACCAATACATGTGTGTTGTGAGGAATGTAATAAGATAATACAGAAGGC CTTGTCATCTACTAAGGATGTAATAATAAAAGGGATGGTCTGTGGCTATAAAGAACAAGGAAAGGATTCTTGTCAA CCCTGGCAGGTGAGCGTGAGGACCAAAGGCAGGCACATCTGCGGCGCGCACCCTGGTCACCGCCACGTGGGTG AATTTCCAGGTGGAAGGTAGGACCAGGTGCTGGGTGACCGGATGGGGCAAAACACCCAGAACGTGGAGAAAAC GGAGATTCTGGGGGCCGCTTGGCCTGTGAATATAATGACACATGGGTCCAGGTAGGGATTGTGAGCTGGGGCA <u> ATCCTCACCCCAGTGTGTGGCCGAACCCCTCTGAGAATCGTGGGAGGAGTGGACGCGGGGAGGAAGGGAGGTGG</u> GACCTTGCCCTTCTCCAGCTCCAACATCCTGTGAATTTTACCTCAAACATCCAGCCTATCTGCATCCTCAGGAG AACACAAGTGTGGGTCTCAGTCCAAAGAGCTTTTGTCCACCCTAAGTTCTCAACAGTTACAACCATTCGAAAT rcgctgtggtcgc

SGPR447\_SEQID\_26

SCAGCCTGCGCCTGAGGAGGCGCCACCGATGTGGAGGGAGCCTGCTCAGCCGCCGCCGCTGGGTGCTCTCGGCTG ATGGGCGCGCGCGCGCGCTGCTGCTGCCGCTGCTGCTCGGCTCGGCTTGGACTCGGGAAGCCGGAGGCCTG

TCCGGAGGGTGATGTCCCACAGTACACCCAGGCCAAACCCCTCCCAGGTGTTGCTGCTCCTTGCCTGCTGTGG CTGTTTGAACAGCCCTCTAGCCGTAGTATGATCTGGGATTCCATGTTTTGTGCTGGTGCTGAGGATGGCAGTGT SAGCTGGGGAATGGACTGCGGTCAACCCAATCGGCCTGGTGTCTACACCAACATCAGTGTGTACTTCCACTGGA <u> GECACACCTCTGCCACCTCCTTACAACCTCCGGGAAGCACAGGTCACCATCTTAAACAACACCAGGTGTAATTAC</u> CCTTGGAACCTGCGGGCCTACAGCAGTCGTTACAAAGTGCAGGACATCATTGTGAACCCTGACGCACTTGGGGT CGCACTGCTTCCAAAAGCACTACTATCCCTCCGAGTGGACGGTCCAGCTGGGCGAGCTGACTTCCAGGCCAACI <u> SACACCTGCAAAGGTGACTCAGGTGGACCCTTGGTCTGTGACAAGGATGGACTGTGGTATCAGGTTGGAATCGT</u> ITTACGCAATGACATTGCCCTGCTGAGACTGGCCTCTTCTGTCACCTACAATGCGTACATCCAGCCCATTTGCAT SGAGTCTTCCACCTTCAACTTCGTGCACCGGCCGGACTGCTGGGTGACCGGCTGGGGGTTAATCAGCCCCAGT

>SGPR432\_1\_SEQID\_27

CCTGGCTGCAGGCTCGAGTTCAGGGGGCCAGCTTTCCTGGCCCAGAGCCCCAGAGACCCCGGAGATGAGTGATGA <u> ATCAGCTGTTCGCGCGTCTTCTCCTCCAGGTGGGGCAGGGGGTTTCGGGCTGGTGGAGCATGTGCTGGGACAGG</u> SCTGCCCACTGCTTCATTGGGCGCCCCAGGCCCCAGAGGAATGGAGCGTAGGGCTGGGGACCAGACCGGAGGAG 4CAGCATCCTCAATCAATCCAACAGCATATTCGGTTGCATCTTCTACACACTACAGCTATTGTTAGGTCTTCAAGC SCTGCACCAGCGACACCTGTCCAACCCGGCCCGGCCTGGGATGCTATGTGGGGGCCCCCAGCCTGGGGTGCA FIGOT GEOCCA GOOT GEACACT GEGA GOO A GOOT GO GO COT COT GOO COT A TOOT GACCACCACOT GOO SECTCAGCGTGCCTGTGGACAGCGTGGCCCCGGCCCCCCAAGCCTCAGGAGGGCAACACACAGTCCCTGGCGA STGGCCCTGGCAGGCCAGTGTGAGGAGGCAAGGAGCCCACATCTGCAGCGGCTCCCTGGTGGCAGACACCTG CTCTGCAGCGTGAGGGACTCAGCCCTGGGGCCGAAGAGGTGGGGGGTGGCTGCCCTGCAGTTGCCCAGGGCCT SAGGCCAGGCTGATGCACCAGGGACAGCTGGCCTGTGGCGGAGCCCTGGTGTCAGAGGAGGCGGTGCTAACT <u> ACGCGCTGCACGTGAAGGCGGCGCGCGCCCGGGACCGGGATTACCGCGCGCTCTGCGACGTGGGCACCGCC</u> SATECTCCTGGGACCCTACGCAATCTGCGCCTGCGTCTCATCAGTCGCCCCACATGTAACTGTATCTACAACCA SATCATCAGCTTTGCATCAAGCTGTGCCCAGGAGGACGCTCCTGTGCTGCTGACCAACACACAGCTGCTCACAGTT 3GTCCTCACTGCTGCCCACTGCTTTGAAAAGGCAGCAGCAACAGAACTGAATTCCTGGTCAGTGGTCCTGGGT

FIG. 1BB

CCGGCGGTCTTCACCGCGCTCCCTGCCTATGAGGACTGGGTCAGCAGTTTGGACTGGCAGGTCTACTTCGCCG CCGGGGGATGGTGTGTACCAGTGCTGTGGGTGAGCTGCCCAGCTGTGAGGGCCCTGTCTGGGGCCACCACTGGTG CATGAGGTGAGGGGCACATGGTTCCTGGCCGGGCTGCACAGCTTCGGAGATGCTTGCCAAGGCCCCGCCAGG 

>SGPR529 SEQID 28

36GATGACCACCTGCTGCTTCTTCAGGGCGAGCAGCTCCGCCGGACGACTCGCTCTGTTGTCCATCCCAAGTAC ATGAGAGCTCCGCACCTCCACCTCTCCGCCGCCTCTGGCGCCCGGGCTCTGGCGAAGCTGCTGCCGCTGCTGCT <u> AGGTTGCTGGCTGGGGCCCCCGGCCCCCGGAGAGTGAAGTACAACAAGGGCCTGACCTGCTCCAGCATCAC</u> CACCAGGGCTCAGGCCCCATCCTGCCAAGGCGAACGGATGAGCACGATCTCATGTTGCTAAAGCTGGCCAGGC SCGTAGTGCCGGGGCCCCGCGTCCGGGCCCTGCAGCTTCCCTACCGCTGTGCTCAGCCCGGAGACCAGTGCC <u> ACCEGEGCCAGGACCCTTGCCAGAGTGACTCTGGAGGCCCCCTGGTCTGTGACGAGGCCCTCCAAGGCATCCT</u> SCGCCCCGTGCGCGCGCGGCTCGCAGCCCTGGCAGGTCTCGCTCTTCAACGGCCTCTCGTTCCACTGCGCGG ATCCTGAGCCCTAAAGAGTGTGAGGTCTTCTACCCTGGCGTGGTCACCAACACAAGATATGTGCTGGACTGG TGGCGCAACTCTGGGCCGCAGAGGCGGCGCTGCTCCCCCAAAACGACACGCGCTTGGACCCCGAAGCCTATG STGTCCTGGTGGACCAGAGTTGGGTGCTGACGGCCGCGCACTGCGGAAACAAGCAGCCACTGTGGGCTCGAGTAG STCGTGGGGTGTTTACCCCTGTGGCTCTGCCCAGCATCCAGCTGTCTACACCCAGATCTGCAAATACATGTCCT SGATCAATAAAGTCATACGCTCCAACTGA

FIG. 1CC

>SGPR428\_1\_SEQID\_29

36TCATGGGAACGAGACATTCAGCAACATCCACTCGGAGAGAAAGCAAGTGCAGAAGGTCATTATTCACAAATA <u> CCAGGAGAGTAGGGGGCCTCTGGTTTGCCAGAAAAAGAACAAAAGCACATGGTACCAGCTGGGTATTATCAG</u> STEGEGETETEGECTETEGCCAGAAGAACATGCCTGGAGTGTACACCGAGTTGTCCAATTATCTGCTTTGGATCG **AAAACGTACAATGTGGCCACAGGCCTGCTTTTCCAAACTCGTCATGGTTACCATTTCATGAACGGCTTCAAGTCC** <u> AGAATGGTGAGTGCCCGTGGCAAGTGAGTATCCAGATGTCACGGAAACACCTCTGTGGAGGCTCAATCTTACAT</u> TACAAACCGCCCCAGCTCGACAGTGACCTCTCTCTGCTTCTACTTGCCACCACCAGTGCAATTCAGCAATTTCAA AATGCCTGTCTGCCTGCAGGAGGAGGAGAGGACCTGGGACTGGTGTTGGATGGCACAGTGGGTAACGACAAT GGGTATGACCAATATGATGACTTAAACATGCACCTGGAAAAGCTGAGAGTGGTGCAGATTAGCCGGAAAGAATG GGTGGTGGGTTCTGACAGCCGCACACTGCTTCCGAAGAACCCTATTAGACATGGCCGTGGTAAATGTCACTGT

AGAGGAAGACTGTGCTGGCAGGGAAGCCGTATAAGTATGAGCCAGACTCTGTGTACGCTTTGCTTCTCTCACCC TGGGCCATCCTGTTACTGTATTTTGTGATGCTTCTATTATCCT

>SGPR425 SEQID 30

GACTATGTCAAAGGGAGTAAAAAGCTAAGGGTAGGGTTGTTGAAGATGAGGAATAAAAGTGGAGGCAAGAACG GAGAGGCGAAGGGTGGGAGAAGAAGAAAAATCTGGCCGGGGTCAGAGGATTGCCGAAGGGAGGCCTTCCT GTTTTGCAGTGTGTCCGACGAATCCAATGATCTCCTTTACCAATACTGCGATGCTGAGTCGGGCTCCACCGGTTC rccacegectetagtegcattctcatttcccctcagcatettctaactecteccactetgttcateategaaag SATCAAGAAAATGCCTGGTGGAATGATCCACTTCTCAGGATTTGATAACGATAGGGCTGATCAGTTGGTCTATCG **GGGGGTCTATCTGCGTCTGAAAGATCCAGACAAAAGAATTGGAAGCGCAAAATCATTGCGGTCTACTCAGGGC** <u> ACCAGTGGGTGGATGTCCACGGGGTTCAGAAGGACTACAACGTTGCTGTTCGCATCACTCCCCTAAAATACGCC</u> TTCTGAATTGGAGGATTATCTTTCCTATGAGACTGTCTTTGAGAATGGCACCCGAACCTTAACCAGGGTGAAAGT TTCAGTGGACCCGGGTCAAGAATACCCACATTCCGAAGGGCTGGGCACGAGGAGGCATGGGGGACGCTACCTT GGCACCGACAGCAGGTTCAGCATCTTGGACAAAAGGTTCTTAACCAATTTCCCTTTCAGCACAGCTGTGAAGCT GATTTTATGTGGCACTTGAGAAAGGTACCCCGGATTGTCAGTGAAAGGACTTTCCATCTCACCAGCCCCGCATT CAGATTTGCCTCTGGATTCACGGGAACGATGCCAATTGTGCTTACGGCTAA

FIG.

1DD

**SGPR548 SEQID 31** 

TGTTGCTGCGCCTAGTCCAGCCCGCACGCCTGAACCCCCAGGTGCGCCCCGCGGTGCTACCCACGCGTTGCCC CCACCCGGGGGGGCCTGTGTGTGTTCTGGCTGGGCCTGGTGTCCCACAACGAGCCTGGGACCGCTGGGAG <u> ATGGGGGACCCAGAAGGAAGCGCAGAGTGGGGTTGGGGGAAGGGGGATACCGGTGGTCAGAAGAAATTTATTAA</u> CAGTGGATGGGATAAGTCTGTGTCTGGAGGGATCCTGGTGGAGGCAGAAGGGTCCTGCCTCACCTGGATTCTC1 GGCAAGTGGCTCTCTACGAGCGTGGACGCTTTAACTGTGGCGCTTCCCTCATCTCCCCACACTGGGTGCTGTCT SCGGCCCACTGCCAAAGCCGCTTCATGAGAGTGCGCCTGGGAGAGCACAACCTGCGCAAGCGCGATGGCCCA CTGGCATCCACAGCAGCCCAGGATGGTGACAAGTTGCTGGAAGGTGACGAGTGTGCACCCCCACTCCCAGCCAT

CCCCCGGTCACAAGTGAGTCTCCCAGATACGTTGCATTGTGCCAACATCAGCATTATCTCGGACACATCTTGTGA TGAGGGTGACTCTGGGGGACCCCTGGTCTGTGGGGGCATCCTGCAGGGCATTGTGTCCTGGGGTGACGTCCCT CAAGAGCTACCCAGGGCGCCTGACAAACACCATGGTGTGTGCAGGCGCGGGGGGGCAGAGGCGCAGAATCCTG TGTGACACACCCACCAAGCCTGGTGTCTATACCAAAGTCTGCCACTACTTGGAGTGGATCAGGGAAACCATGAA GAGGAACTGA

# >SGPR396 SEQID 32

CACCTTCACTTCTGCCATCCTGCCTATTTGCTTGCCCAGTGTCACAAAGCAGTTGGCAATTCCACCCTTTTGTTGG ATGGGCCCTGCTGGCTGTGCCTTCACGCTGCTCCTTCTGCTGGGGATCTCAGTGTGTGGGGCAACCTGTATACTC CAGCCGCGTTGTAGGTGGCCAGGATGCTGCTGCAGGGCGCTGGCCTTGGCAGGTCAGCCTACACTTTGACCAC TGTCGTGTCACATTGATGGTGTATGGATCCAGACAGGAGTAGTAAGCTGGGGGATTAGAATGTGGTAAATCTCTTC CATTATTGACCGCCAGGCTTGTGAACAGCTCTACAATCCCATCGGTATCTTCTTGCCAGCACTGGAGCCAGTCAT STCTGACTTCTTGTTCCCTATTGTCCTACTCTCTCTGGCTCTCCTGCGTCCCTCCTGTGCCTTTGGACCTAACACT GTCCAAAATCGTCATCCATCCCAAGTACCAAGATACAACGGCAGACGTCGCCTTGTTGAAACTGTCCTCTCAAGT GTGACCGGATGGGGAAAAGTTAAGGAAAGTTCAGATAGAGATTACCATTCTGCCCTTCAGGAAGCAGAAGTACC CAAGGAAGACAAGATTTGTGCTGGTGATACTCAAAACATGAAGGATAGTTGCAAGGGTGATTCTGGAGGGCCTC AACTITATCTGTGGAGGTTCCCTCGTCAGTGAGAGGTTGATACTGACAGCAGCACACTGCATACAACCGACCTG GACTACTTTTCATATACTGTGGGTAGGATCGATTACAGTAGGTGACTCAAGGAAACGTGTGAAGTACTACGT CTGGAGTCTACACCAATGTAATCTACCAAAAATGGATTAATGCCACTATTTCAAGAGCCAACAATCTAGACT rcccaggggcagataa

FIG. 1EE

# >SGPR426 SEQID 33

STIGITGAGGATGATAAGTCTTTCTATTACCTTGCCTCTTTTAAAGTCACAAATATCAAATATAAAGAAAATTATGG <u> ACTCAGTACGGCTAGCTCTTTTCACATTAGCAATTGTAGCAATCATAGGAATTGCAATTGGTATTGTTACTCATTTT</u> TGAAGACCAAACAATTGTCTTTGACCATAAACAAACCATCATTTAGACTCACACGCTGTGGAATAAGGATGACAT ICTGTAGGCGGTCGATTTATCAAATCTCATGTTATCAAATTAAGTCCAGATGAACAAGGTGTGGATATTCTTATAG CATAAGATCTTCAAGAGAGTTTATAGAAAGGAGTCATCAGATTGAAAGAATGATGTCTAGGATATTTCGACATTCT 

TATAACACCACCGCAGTGAAACGAAATGTGAGGAAAATTATTCTTCATGAGAATTACCATAGAGAAACAAATGAA CATCTATAAAGTTGCCACCTAAAACAAGTGTGTTCGTCACAGGATTTGGATCCATTGTAGATGATGGACCTATACA AAATACACTTCGGCAAGCCAGAGTGGAAACCATAAGCACTGATGTGTGTAACAGAAAGGATGTGTATGATGGCCT ICTEGTTTATGATAATCATGACATCTGGTACATTGTAGGTATAGTAAGTTGGGGACAATCATGTGCACTTCCCAAA AATGACATTGCTTTGGTTCAGCTCTCTACTGGAGTTGAGTTTTCAAATATAGTCCAGAGAGTTTGCCTCCCAGACT A T G G C T G C T C A C T G C T T T T G G A A A A A A G A C C C A A T G G A T T G C T A C T T T G G T G C A A C <u> AAACCTGGAGTCTACACCAGAGTAACTAAGTATCGAGATTGGATTGCCTCAAAGACTGGTATĠTAG</u>

>SGPR552\_SEQID\_34

**AAGCATGTCTTCCAGATGTTAATTATATATTCCTATACAATTCAGAAGCAGTGGTTACTGCATGGGGATCATTTAAA** <u>AGAATAGCAGAGGGTCTGGATGCTGAGGAAGGAGTGGCCCTGGCAAGCTAGCCTTCCACAGAACAATGTCTA</u> CCAAAGAATGGAATGTTATTTTAAGTAACCCACAAACACAGTCAAATATCAAGAATGTTATAATTCAAGAAACTAC AGGGTGACTCTGGTGGACCACTGGTTGGTACAGATTCTAAAGGCATCCTTGCTAAAGGTTCCCTGCTGGTATTGA CCGACGCGGAGCCACATGGCTTAGTAACAGCTGGCTTATCACTGCTGCTCACTGCTTCATAAGGGTCCATGATC CCTTTACGAACAACTTCTAATGTACTCCACAAGGGATTAGTGAAGATTATAGATAATAGGACCTGCAACAATGGG **AAGCTGGAGTAAATGAACGTGCTCTTCCAAACAAGCCTAGTGTCTACACTCAAGTGACATACTAT** 

FIG. 1FF

>SGPR405 SEQID\_35

CCCTCACAGGTCCGGGCAGCCTTCCTAGAGGAGGTGGCATTGAGGTGGGCATGGAGTTTCCGGGATGCAGCGG GACCCAAGCGACGGCAGCAGGGGTCAGCAGAGGGGCCTGCGGCGGGGGTGGACGCTGGAGCAGGAGACCAGG <u> ATGGTCAĞCAAGĞGGGGAGTTGCTGCAGAGCCAGAGCCACACTATTGTGAGGGACAGTGAAAGAGGGCCCCAACA</u> GTTTAGAGTGCAGGTGGGGCAGCTGAGGCTCTATGAGGACGACCAGCGGACGAAGGTGGTTGAGATCGTCCGT CACCCCCAGTACAACGAGAGCCTGTCTGCCCAGGGCGGTGCGGACATCGCCCTGCTGAAGCTGGAGGCCCCG TCCCAATAGACAGCAAACACCTGAAACCGGTGATCAGCAGCTTCCCGGTAAGATCTCAGGAGCTGGGCGAGGG GGCTGGAGCAGGCACACTAAGAGGCAAAATGGCAGAGTTTAACTGGTCTATGGCCTTCAAGGGACCTGCGGCT GGTCATGAAGAGCGCCTCAACTCTGTGTCCAGCAGGGCCAAGAAGGGCCATTGGCTGGGATGTCGCTGCTGCTT

GCAGGTGAAGGTCCTCACACTGAGCAATGCAGACTGTGAGCGGCAGACCTACGATGCTTTTCCTGGTGCTGGAG CGGCCCAGCATTGGCGTCTACACACGCCCAGAGACCAGCTGGCAGGGTGCCAACCATGCAGACGCCCAGAGAC CCTGCTACCCAATGGCTGTCCCCAGCACCCTTACCCACGTCACCTGCCACCCGGCGGCCATCCCCAGGCCCTTC GCTGGGTGACCGGCTGGGGTGTCATTGGACGTGGAGAACTACTGCCCTGGCCCCTCAGCTTGTGGGAGGCGAC A C C CACATCACCT G C TACACGAT G G C CAT C C C C G G C C T C C A C C C C C C C T G C T A C C G G C C A T C C C C GTGGTCGTGGCAGTGGGTGCTGACCGCCGCTCACTGCATTTTCCGGAAGGACACCGACCCGTCCACCTACCGG TCACCCACGTCACCTGCTACCCGACGGCCATCCCCAGGCCCTTCACCCATGTCACCTGCTACCTGATGGCTGTC CCCAGCACCCTCACCCACGTCACCTGCTACCCGACGGCCGTCCCCAGGCCCTTCACCCATGTCACCTGCTACCT AGCACCCTCACCCACGTCACCTGCTACACGATGGCCGTCCCCAGGCCCATCACCCATGTCACCTGCTACACGAT GEGECCCCTCCTGTGCAGGCGGAATTGCACCTGGGTCCAGGTGGAGGTGGTGAGCTGGGGCAAACTCTGCG TGATGGGATCAGGAGCGCCGCTGCCCCCGGCCCCCGACCTGCAAGAGGCCGAGGTCCCCATCATGAGGACCC GTGTGCAGGGAGGAAGGGGCAGGGTTCCTGCCAGGCCGCTCTGAGGACGGAGGACCTCACCCCAACCACACC CTACTCTGTCTTCCTGGGGGGGAGACATCGCCCTGCTGAAGCTGGCCACCAGTTCCCTGGAGTTCACTGACA 26GCCCCCTGGTCTGCAAGAAGAAGGGTACCTGGCTCCAGGCGGGAGTAGTGAGCTGGGGATTTTACAGTGAT <u> GATGGCTGTCCCCAGCACCCTCACCCACATCACCTGCTACATGATGGCCGTCCCCAGGCCCTTTACCCACATCA</u> GAGCTTGCGAGAGGATGTATCACAAAGGCCCCACTGCCCACGGCCAGGGTCACCATCATCAAGGCTGCCATGCC <u> 2AGCTGGCAGGGTGCCAACCATGCAGAGGCCCAGAGACATGGGCCAGGGCCAGGAGTGGGTCTGCAGGCCCT</u> TTGAGCGGCTCATCAAGGACGACATGCTGTGTGCCGGGGACGGGAACCACGGCTCCTGGCCAGGCGACAACG AGCTCAGACCCCTGCTGTGGTCCGAAGATTTGTGCTCCCCCCCAAATCCAGATGTTGAAGCCCTAACTCCAGTG CAACACGGAGGTGTCTCCACGTGCAGACCCCAGGCTGAGCCAGGCGGAGGAGATCTGGCCAGAGTGGGCTTG <u> ACTGGGAGTCCAGCTCCCGTCCCCGAGAATGACCTGGTGGGCCATTGTGGGGGGCCACACACCCCCAGGGGAA</u> **ATTCACACCAGGGATGTGTATCTGTACGGGGGCCGGGGGCTGCTGAATGTCAGCCAGATCGTCGTCGTCCACCCAA** GTGCCGCTGTCTGAGCTCATCCACCCGGTCTCGCTCCCGTCTGCCTCCCTGGACGTGCCCTCGGGGAAGACCT

SGPR485 1 SEQID 36

4GGGTGACAGTGGGGGCCTCTGGTCTGCACCCCAGAGCCTGGTGAGAAGTGGTACCAGGTGGGCATCATCAG **AACAAGTGGTGGATTCTCACTGCGGCTCACTGCTTATATTCCGAGGAGCTGTTTCCAGAAGAACTGAGTGTCGTG** CTGGGGAAAGAGCTGTGGAGAGAAGAACACCCCAGGGATATACACCTCGTTGGTGAACTACAACCTCTGGATCG 26GAGGTGGGTGAGTTTCCGTGGCAGGTGAGTATTCAGGTAAGAAGTGAACCTTTCTGTGGCGGCTCCATCCTC AGAAAGTGACCCAGCTAGAGGGCAGGCCCTTCAATGCAGAGAAAAGGAGGACTTCTGTCAAACAGAAACCTATG 4GTCAGTGAATGTGGTGACAGATCTATTTTCGAGGGAAGAACTCGGTATTCCAGAATCACAGGGGGGGATGGAGG SCCCATCTGCCTCCCCACGCAGCCCGGCCCTGCCACATGGCGCGAATGCTGGGTGGCAGGTTGGGGGCAGAC SECTCCCCAGTCTCGGGAGTCCCAGAGCCAGGCCAGGCCCCAGATCCTGGCTCCTGCTCTGTCCCTGTCCATG CAATGCTGCTGACAAAACTCTGTGAAAACGGATCTGATGAAAGCGCCAATGGTCATCATGGACTGGGAGGAGT GTTCAAAGATGTTTCCAAAACTTACCAAAAATATGCTGTGTGCCGGATACAAGAATGAGAGCTATGATGCCTGCA 4GAGGCTGGAGTGGCTATCCTAGGCAGGGCTAGGGGAGCCCACCGCCCTCAGCCCCTCATCCCCCAGCCC 4AGAGAGCCAACATGGACAATGACATTGCCTTGCTGCTGCTTCGCCCATCAAGCTCGATGACCTGAAGGT A T G C T C T G T T C T C A G T G C T G C T C C T G G T C A C G A C T C A G C T C G G T C C A C G A C T C C T C C C STGGGGACCAACGACTTAACTAGCCCATCCATGGAAATAAAGGAGGTCGCCAGCATCATTCTTCACAAAGACTT IGTTGTTCAGAGCTATTTTGTACTGA

>SGPR534 SEQID 37

FIG. 1HH

ATCCAGGTCCTGAAGATCGCCAAGGTCTTCAAGAACCCCAAGTTCAGCATTCTGACCGTGAACAATGACATCACC <u>GACGTGATCTGTGCCGGGGCCAGTGGCGTCTCCTCCTGCATGGGTGACTCTGGAGGCCCCCTGGTCTGCC</u> ACTTCCCCGCGGGGACACTGTGTGCCACCACAGGCTGGGGCAAGACCAAGACGACAACAACAAGACCCTGA SAAGCTGCAGCAGGCAGCCCTGCCCTCCTGTCCAATGCCGAATGCAAGAAGTCCTGGGGCAGGAGGATCACC SCCCACTECEGGGTCAGGACCTCCGACGTGGTCGTGGCTGGGGAGTTTGACCAGGGCTCTGACGAGGAGAAC STCCCTGCAGGACAAAACCGGCTTCCACTTCTGCGGGGGCTCCCTCATCAGCGAGGACTGGGTGGTCACCGCT CCCTGTGCTCAGCGGCCTGTCCAGGATCGTGAATGGGGAGGACGCCGTCCCCGGCTCCTGGCCCTGGCAGGT 4GAAGGACGGAGCCTGGACCCTGGTGGGCATTGTGTCCTGGGGCAGCCGCACCTGCTCTACCACCACGCCCG STGTGTACGCCCGTGTCACCAAGCTCATACCCTGGGTGCAGAAGATCCTGGCCGCCAACTGA

FIG. 111 SCTGCGGGAACAGCCAGTGTGTGACCAAGGTGAACCCGGAGTGTGACGACCAGGAGGACTGCTCCGATGGGTC SAGCCCTCTGGCCGGTTCTTTCTGGCTGGCATCGTGAGCTGGGGAATCGGGTGTGCGGAAGCCCGGCGTCCAG 4GCGAGGGATCCGGGCAAGGCTGCGGGAGCACGGCATCTCCCTGGCTGCCTATGGCACAATTGTGTCGGCTGA A T CAAC G C C A G G T G G T G T G C T G C T G C T T C A A T G A G T T C C A G A C C C G A G T G G G T G G C T SCCCACCATGGCTCCTGCCCCTGCCGCCCCCAGCACAGCCTGGCCCACCAGTCCTGAGAGCCCTGTGGTCAGC CGACGAGGCGCACTGCGAGTGTGGCTTGCAGCCTGCCTGGAGGATGGCCGGCAGGATCGTGGGCGGCGGCATGGA 4GCATCCCCGGGGGAGTTTCCGTGGCAAGCCAGCCTTCGAGAGAACAAGGAGCACTTCTGTGGGGGCCGCCATC SCTGGACTTCGACCTGGCTGTCCTGGAGCTGGCCAGCCCCTGGCCTTCAACAAATACATCCAGCCTGTCTGCC SGGTCTATGCCCGAGTCACCAGGCTACGTGACTGGATCCTGGAGGCCACCACCAAAGCCAGCATGCCTCTGGC GCCCCTGGCCATCCAGAAGTTCCCTGTGGGCCGGAAGTGCATGATCTCCGGATGGGGAAATACGCAGGAAGG **AAATGCCACCAAGCCCGAGCTCCTGCAGAAGGCGTCCGTGGGCATCATAGACCAGAAAACCTGTAGTGTGTCTCT** CETGGGTGCGACCTACCTCAGCGGCTCGGAGGCCAGCACCGTGCGGGCCCCAGGTGGTCCAGATCGTCAAGCA SCCCCTGTACAACGCGGACACGGCCGACTTTGACGTGGCTGTGCTGGAGCTGACCAGCCCTCTGCCTTTCGGC STGTGCGCTGGCTACCTGGACGGGAAGGTGGACTCCTGCCAGGGTGACTCAGGAGGACCCCTGGTCTGCGAG ACCCCCACCAAATCGATGCAGGCCCTCAGTACCGTGCCTCTTGACTGGGTCACCGTTCCTAAGCTACAAGAATG 36GCCTGGGCGGGAGCCCGGTGAAGATCGGGCTGCGGCGGGTAGTGCTGCACCCCCCTCTACAACCCTGGCAT IGTCTGCCGCCCACTGCTTCAACCACACGAAGGTGGAGCAGGTTCGGGCCCCACCTGGGCACTGCGTCCCTCCT SAAAGCCACTGTGGAGCTGCTGGACCAGGCACTGTGTGCCAGCTTGTACGGCCATTCACTGACAGGATG 'GGGGCCAGGCCTGCAATGGAGAAGCCCACCCGGGTCGTGGGCGGGTTCGGAGCTGCCTCCGGGGGGGTGC GCTGGGGCTACCTCAAGGAGGACTTCCGTAAGCATCTTCCTCGGCCTGCAATGGTCAAGCCAGAGGTGCTGCA SCTCACAGGGAGACATAAGGGACCCTTGGCAGAAAGAGACTTCAAATCAGGCCGCTGTCCAGGGAACTCCTTT

SACACCATCTTTAAGAAATGTAACACTCAGCCCTTCCTCTACACAGGGCTTCCACGTGGACCACACGGCCGAG

STGCGGGGGAATCCGGTGGACCAGCAGTTTGCGGCGGGAGACCTCGGACTATCACCGCACGCTGACGCCCACC

SAAGCAAGACCACGAACCCACCAGAAGAAGAAATTCCGAACATCCGAACATCAGCAAGAAGAACAACTCCGGA AATGAACAACTCCAGACACGCTGCCTTAAGAGCTGCAACACTCCCTGGGAAGGTCTACAGCGTCACTCCTGAAG

SCTGTCGAGCGGCCAGCATTGGCGTGGTGGCCACCAGCCTTGTCGTCCTCACCCTGGGAGTCCTTTTGGGAGG

<u> ATGGAGCCCACTGTGGCTGACGTACACCTCGTGCCCAGGACAACCAAGGAAGTCCCCGCTCTGGATGCCGCG</u>

>SGPR390 SEQID 38

STGGAGGCACTGCTGCACTTTCTGCTGCGACCCCTCCAGACGCTGAGCCTGGGCCTGGAGGAGGAGGAGTTGC

AGCGAGCAGACCTGCCGCCGCTTCTACCCAGTGCAGATCAGCAGCCGCATGCTGTGTGCCGGCTTCCCGCAGG TCTGGGGGCCCCCTGGCGAGGAGGCCCCTGGCGTGTTTATCTGGCAGGGATCGTGAGCTGGGGTTT ACAACTTCTCCCTCACAGACCGCATGATCTGCGCAGGCTTCCTGGAAGGCAAAGTCGACTCCTGCCAGGGTGAC CAGCGGCCCGTGGGGAGTGGCCGTGGCAGGTGAGCCTGTGGCTGCGGCGCCGGGAACACCGTTGCGGGGCC GTECTEGTEGCAGAGAGGTGCTGCTGTCGGCGCGCGCACTGCTTCGACGTCTACGGGGACCCCAAGCAGTGG CCACCACACACACCACCCAGCTACCAGACTGTGGCCTGGCGCCGCCGCGCGCTCACCAGGATTGTGGGCGGCGCGCG GTCCTCCCAGCCCCTTCCCATGTCTCCCCCCTCGACCACAAGGATGCTGGCCACCACCAGCCCCAGGACGACA CCTGCCAACTCAACCTTATCTGCCGTGAGCACCACTGCTAGGGGACAGACGCCATTTCCAGACGCCCGGAGG GCTGCGCTCAGGTTAAGAAGCCGGGCGTGTACACGCGCATCACCAGGCTAAAGGGCTGGATCCTGGAGATCA1 IGTGAGAGGCTGGATAGGACAGCACATCCAGGAGTGA

>SGPR521\_SEQID\_39

GGCTGTCATCCATGGTGAAGAAAGTCAGGCTGCCCTCCCGCTGCGAACCCCCTGGAACCACCTGTACTGTCTCC ATGAGTACACCGTGCACCTGGGCAGTGATACGCTGGGCGACAGGAGGGCTCAGAGGATCAAGGCCTCGAAGTC GGCTGGGGCACTACCACGAGCCCAGATGTGACCTTTCCCTCTGACCTCATGTGCGTGGATGTCAAGCTCATCTC GGCAATCAGCTCCACTGCGGAGGCGTCCTGGTCAATGAGCGCTGGGTGCTCACTGCCGCCCACTGCAAGATGA CCCCCAGGACTGCACGAAGGTTTACAAGGACTTACTGGAAAATTCCATGCTGTGCGCTGGCATCCCCGACTCCA AGAAAAACGCCTGCAATGGTGACTCAGGGGGACCGTTGGTGTGCAGAGGTACCCTGCAAGGTCTGGTGTCCTG ATTCCGCCACCCGGCTACTCCACACACAGACCCATGTTAATGACCTCATGCTCGTGAAGCTCAATAGCCAGGCCA GGGAACTTTCCCTTGCGGCCAACCCAATGACCCAGGAGTCTACACTCAAGTGTGCAAGTTCACCAAGTGGATAA ATGGCAAGATCCCTTCTCCTGCCCCTGCAGATCCTACTGCTATCCTTAGCCTTGGAAACTGCAGGAGAAGA ATGACACCATGAAAAAGCATCGCTAA

>SGPR530\_1\_SEQID\_40

GCCTGGGTCCTGGTGGGGCTGGCCAGCTGGGCCTGGACTGCCGGCATCCTGCCTACCCCAGCATCTTCACCA GGGTCACCTACTTCATCAACTGGATTGACGAAATCATGAGGCTCACTCCTCTTTCTGACCCCGCGCTGGCTCCTC SAAGACCATAAGAGGGTGCAACTGTCACCACCCTTCTATCTCCAGGAGGGCAAGGTGGGCCTCATTGAGAACAC GGTATCAACTACCCACTGCTTTCTCAACAAATCCCAGGCCCCGAAGAACTATCAGGTTCTGTTGGGAAACATCCA GGGGGTACTTCTCGACAGGAAAGTCCATCTGCAAAGGCGATTCTGGGGGGCCTCTAGTCTGCTACCTCCCAGT <u> GTCTCCACAGTGTGTGGGAAGCCTAAGGTGGTGGGGAAGATCTATGGTGGCCGGGACGCAGCAGCTGGCCAGT</u> ACTGTATCATCAAACCCAGCACACCCAGAAGATGTCTGTGCACCGGATCATCACCCATCCAGACTTTGAGAAGCT <u> ACTCTGTAATACCTTATATGGGCAAAGAACTGCAAAGGCGAGACCTAAGCTTTGCACGAGGAGATGCTGTGTGG</u> SCACCCCTTTGGGAGTGACATTGCCATGTTGCAGCTGCACCTGCCTATGAACTTCACTTCCTACATTGTCCCTGT CTGCCTCCCATCCCGGGACATGCAGCTGCCCAGTAACGTGTCCTGTTGGATAACCGGCTGGGGAATGCTCACC SGCCATGGCAGGCCAGCCTGCTCTACTGGGGCTCGCACCTCTGTGGAGCTGTCCTCATCGACTCCTGCTGGCT

>SGPR520 SEQID 41

1KK

<u> ATGCTGCTGCTGCTGCTGCTGCTACCCCTCCCAAGCTCATGGTTTGCCCCACGGGCACCCACTGTACACACG</u> CCTGCCCCCCAGCGCCCTGCAAGTCTTCACTCTCCTTTGGGGGCCAGAGACTGTGTTGGGCCGCAACCTAGACT TGGGGGGCAGCGCGGCGCCCGGGGCCTGGCCCTGGCTGGTGAGGCTGCAGCTCGGCGGGGGGAGCTTTG CCTGCCGGAACCGCCTGCGCCATCGCGGGCTGGGGCGCCCTCTTCGAAGACGGGCCTGAGGCTGAAGCAGTG SCCCGGCTCTGCGCCTTCTATGCCCGCCTGTGCCCGGGGTCCCAGGGCGCCCTGTGCGCGCCTTGGGGGCGCCTGGCGCCAG ACGTTTGTGAAGGGCCGTGCGGCGAGAGGCGTCCGAGCACTGCCAATGTGACGCGGGCCCACGGCCGCATCG ACCATECTCTGCGCCGGGTACCTGGCGGGGGGGGGTTGACTCGTGCCAGGGTGACTCGGGAGGCCCCTGACC TOCTOCAGOCGOGAGOCCAGOTGCAGGGAGOTTCTGGCCTGGGACCCCCCCCAGGAGGTGCAGGCAGACGCC STGCCAGTGAACCGCATCCTGCCCCACCCCAAGTTTGACCCGCGGACCTTCCACAACGACCTGGCCTGGTGC <u> AGAGAGGCCCGTGTTCCCCTGCTCAGCACCGACACCTGCCGAAGAGCCCTGGGGGCCCGGGCTGCGCCCAGC</u> CCAGGGAAGCCCGGGGTCTACACCCGCGTGGCAGTGTTCAAGGACTGGCTCCAGGAGCAGATGAGCGCAGCC TGCGGCGCGCGTCCTGGTAGCGGCCTCCTGGGTGCTCACGGCAGCGCACTGCTTTGTAGGCTGCCGCTCGACC

GCGCAGGAGCTGCTCGGGCCTCGTCCGGGACTGCGGCCCTGGCCCCCGCCCTGGCTCTCCCCGGCTCTCCCAGCG CTCAGGGAGTCTCCTCTGCACCCCGGCCCGGGAGCTGCGGCTTCACTCAGGATCGCGGGCTGCAGGCACTCGGT rcccaagcggaggccggagccgcgcgcgagaagccaacggctgccctgggctggagccctgcgagagt GGTGCAGGCCCTGCAGGCCTTCCGCGTGGCTGCCCTGGCAGAGGGGGAGCCCGAGGGACCCTGGATGTG FGECTGCCCTGCAGGGGCCCCATGCCTGGATCCTGCAGGTCCCCTCGGAGCACCTGGCCATGAACTTTCATGA SCGGCATGTGGCCTTCAGCGGCCTGGTGGGCCTGGAGCCGGCCACACTGGCTCGCAGCCTCCCCCGGCTGCT GGTCCTGGCAGATCTGGGCTCCAAGACACTGACCGGGCTTTTCAGAGCCTGGGTGCGGGCAGGCTTGGGGGG

>SGPR455\_SEQID\_42

TGTGGAATGTTGACCATCACTTCAATATTCAGTTCTAGTAACATGACGGTGATATACTTTAAAAGTGATGGTAAAA STGAGAAGGGCCAAACACATAATAGTGCATGAAGACTTTAACACACTAAGTTATGACTCTGACATTGCCCTAATAC AATATTCAAATGTCCTACAAGAAATGGAACTTCCCATCATGGATGACAGAGCGTGTAATACTGTGCTCAAGAGCAT AAATTTAGTCCAAACTGTATTTATGATGCTGTTGTGATTTACGGTGATTCTGAAGAAAAGCACAAGTTAGCTAAACT CCCCAAAACAATCCTGTATCTACCGTAAAAGCTATTCTGCATGATGTCTGTGGCATCCCTCCATTTAGTCCCAG TGAAGAATAATĆCACTCTCCTGGACTATTATTGCTGGGGACCATGACAGAAACCTGAAGGAATCAACAGAGCAG ATTTACAAGTTTAGACATGGAAAAGCAAGTTGGATGTGATCATGACTATGTATCTTTACGATCAAGCAGTGGAGTG 4GGTTGTGGGAGTCTGGCTATATTGGTAGAAGAAGGGACAAATCACTCTGCCAAGTATCCTGATTTGTATCCCAG : AACACACAAGGTGTCATTGGTTCATTTGTGCTCCAGAGAAGCACATTATAAAAGTTGACATTTGAGGACTTTGCTGTC ATGAGTCCTGATATTGCACTGCTGTATCTAAAACACAAAGTCAAGTTTGGAAATGCTGTTCAGCCAATCTGTCTTC STGACAGCGATGATAAAGTTGAACCAGGAATTCTTTGCTTATCCAGTGGATGGGGCAAGATTTCCAAAACATCAG <u> GAACCTCCCTCCCTGGGAAGGACCATGCTGTGTGCTGGCTTCCCTGATTGGGGAATGGACGCCTGCCAGGGG</u> SACTCTGGAGGACCACTGGTTTGTAGAAGAGGTGGTGGAATCTGGATTCTTGCTGGGATAACTTCCTGGGTAGC TGGTTGTGCTGGAGGTTCAGTTCCCGTAAGAACAACCATGTGAAGGCATCACTTGGCATTTTCTCCAAAGTGTC GAGTTGATGGATTTTATCACTCAAAACCTGTTCACAGGTTTGGATCGGGGCCAACCCCTCTCAAAAGTGGGCTC 'GGCTTTCCAGAAGAATCGCAGGAGGGGAAGAAGCCTGCCCCCACTGTTGGCCATGGCAGGTGGGTTGAGGT **AAGGTATATAACAAAGGCCCTGAGTTCTGTCCAAGAAGTGAATGGAAGCCAGAGAGATAAAATAATCCTGATAAA** 1TCGTTTACAAGGCTTCAAGGCCAGATTTACCATTTTGCCCTCAGAGTCTTTAAACAAATTTGAACCAAAGTTACC TCTAGGCGATTACCAATGTGGAGGTGCCATCATCAACCCAGTGTGGATTCTGACCGCAGCCCACTGTGTGCAA STITCTGATACAGAAGACAGTGGCAGTGGCTTTGAGCTTACCGTTACTGCTGTACAGAAGTCAGAAGCAGGGTC CTTTTTAGTAAGGTCTGTGGAAAAATATTGCCTTCACCATTGCTGGCAGAGACCAGTGAGGCCATGGTTCCATTT

FIG. 1LL

ATGCAGATCCCAAGAAATTCTCACAGAACTACTATGGGTTGCCAATGGAGATTAGTAGCCCCTTTAAATCACATCA 4GACTTTCCCAAAGTTCAAACAGAGAGCACTTGGTCCCTTGTGAGGATGTTCTTCTGACCAAGCCAGAAGGGATC ATTTGGACCAGGAAAAAAATTAATAGGTAGAATGTTGATGAGCACTGAGCTTTCTTGGTTCCTAAGCCAATTCAGC 36GAGCTGGCTGTGCCAGCCATGGAAGCCGGGTGTATTTGCCAGAGTGATGATCTTCTTGGACTGGATCCATAT SGTGGATTACATGGAAGAAGACTTTACTCAATGACTTTCATGAGTCCTGGACCGCTGGTGAGGGTGACATTCCAT STGCCCATCCAGGAGGGATCACAGAGAAGATGATCTGTGCTGGCTTTGCAGCATCTGGAGAGAAAGATTTCTGC AGTTGTTGCTCTGAAGCAGAGGCTAGAAAGCCTAGAGGCTTTTTCCCACACCACGGGTATCTACTGGATTATAGA GCCCTTGTACGAGGTGCATTTGGTATAAGCTATATTGTCTTGAAAGTCCTAGGTCCAAAGGACAGTAAAATAACC CAGGGAGACTCTGGTGGGCCACTAGTATGTAGACATGAAAATGGTCCCTTTGTCCTCTATGGCATTGTCAGCTG STGGCCTAGCAAGTCGCCTACAGCAGATTCAAGTGCATGTGTTAGAAAGAGAGGGTCTGTGAACACACTTACTAT IACAGAACACCTGTTACCATGCACTGCCTCATGAGGTTGTTTTGAGAATTAAATAA

GCTCATCAAGCTGGCTAAGCCTGCCATGCTCAATCCCAAAGTCCAGCCCCTTACCCTCGCCACCACCAATGTCA 36CCAGGCACTGTCTGTCTACTCTCAGGTTTGGACTGGAGCCAAGAAAAACAGTGGGCTTTGGCAGCTGGAGCCA CTGCTCCCTATTTGGTGTACCTCAAGTCTCACTTCAACCCCTGTGTGGGCGTCCTCATCAAACCCAGCTGGGTGC FGGCCCCAGCTCACTGCTATTTACCAAATCTGAAAGTGATGCTGGGAAATTTCAAGAGCAGAGTCAGAGACGGTA CTGAACAGACAATTAACCCCATTCAGATCGTCCGCTACTGGAACTACAGTCATAGCGCCCCACAGGATGACCTCA ACACCCTGACTTGCGGCAGAACCTGGAGGCCCCCGTGATGTCTGATCGAGAATGCCAAAAAAACAGGAACAAGGAA CCAGGCCATCTGACTCTGCACAGAGGCCCCAGCCATTCCTGATTGGCAGAGACACAATTCACATGAACAAGGCCG AAAGCCACAGGAATTCCTTATGTGTGAAATTTGTGAAAGTATTCAGCCGAATTTTTGGGGAGGTGGCCGTTGCTA STGTCATCTGCAAAGACAAGCTCCAGGGAATCGAGGTGGGGCACTTCATGGGAGGGGGGCGTCGGCATCTACAC CAATGTTTACAAATATGTATCCTGGATTGAGAACACTGCTAAGGACAAGTG/

>SGPR559 SEQID 44

SECCCCTETEGATCATCACTECTECACACTGTTTATGACTTGTACCTCCCCAAGTCATGGACCATCCAGGTGG STCTAGTTTCCCTGTTGGACAATCCAGCCCCATCCCACTTGGTGGAGAAGATTGTCTACCACAGGAAGTACAAGC <u> AAGTACAGATGTCGCTCATCCTTTAAGTGTATCGAGCTGATAGCTCGATGTGACGGAGTCTCGGATTGCAAAGAC</u> SGAAGACCATGTGCTCCGATGACTGGAAGGGTCACTACGCAAATGTTGCCTGTGCCCAACTGGGTTTCCCAAGC 
 FATGTGAGTTCAGATAACCTCAGAGTGAGCTCGCTGGAGGGGCAGTTCCGGGAGGAGTTTGTGTCCATCGATCA
NGGATGGAGCAGGTGACGCCTCCCCTGTCCTGAACCACGCGGCCGTCCCTTTGATTTCCAACAAGATCTGCAAC SACAGGGACGTGTACGGTGGCATCATCTCCCCCTCCATGCTCTGCGCGGGCTACCTGACGGGTGGCGTGGACA SCTGCCAGGGGGACAGCGGGGGGCCCCTGGTGTGTCAAGAGAGGAGGCTGTGGAAGTTAGTGGGAGCGACCA <u> ATCATCGTCATTGGGATCATTGCATTGATATTAGCACTGGCCATTGGTCTGGGCATCCACTTCGACTGCTCAGGG</u> GGGGAGGACGAGTACCGCTGTGTCCGGGTGGTGGTCAGAATGCCGTGCTCCAGGTGTTCACAGCTGCTTCGT STGTGCCTGCCCAACTCTGAAGAGAACTTCCCCGATGGAAAAGTGTGCTGGACGTCAGGATGGGGGGCCACAG TECTCTCGCAGTGGCCCTGGCCAGCCTTCAGTTCCAGGGCTACCACCTGTGCGGGGGCTCTGTCATCA CAAAGAGGCTGGGCAATGACATCGCCCTTATGAAGCTGGCCGGGCCACTCACGTTCAATGAAATGATCCAGCCT ATGGGGGAAAATGATCCGCCTGCTGTTGAAGCCCCCTTCTCATTCCGATCGCTTTTTGGCCTTGATGATTTGAAA <u> ATAAGTCCTGTTGCACCAGATGCAGATGCTGTTGCTGCACAGATCCTGTCACTGCTGCCATTGAAGTTTTTCCA</u> CCACGAGCAGATGGAGAGAGGACCTAAAAAACCTGA

>SGPR567 1 SEQID 45

1NN

STAGCAGCAACTGGAATGACTCCTACTCAGAGAGACCTGCCAGCAGCTGGGTTTCGAGAGTGCTCACCGGACA CTCCAGCTGGGACACCTCCAGGCCGGGCATCTCCAGCCCAGGCATCTCCAGCCCAGGCATCTCCAGCTGGGAC CAGAGTGTACCTTGTTAGAGCAACACCAGTGGGGGCTGTACCCATCCGATCATCTCCTGCCAGGTCAGCACCAG CTGGCATCACTTTCCAGGTCCTCATCCGGCAGGTCATCATCCGCCAGGTCAGCCTCGGTGACAACCTCCCCAAC GCTGCGTGAGGTTTGACTGGGACAAGTCTCTGCTTAAAATCTACTCTGGGTCCTCCCATCAGTGGCTTCCCATC 36CATCTCCAGCCCAGGCATCTCCAGCCCAGGCATCTCCAGCCCAGGCATCTCCAGCCCGGGCATCTCCGGCT SEGAGAGCTETCCCAAGCACGCTGTTCGCTGTGACGGGGTGGTGGACTGCAAGCTGAAGAGTGACGAGCTGG CAACCAGGGCCACCAGGGAGAGCCCAGTCCAGTTCTGGCAGGGCCACACAGGGATCAGGTACAAGGAGCAGA ACCTCCGGGCCGGGCATCTCCAGCCCAGGCATCTCCAGCTGGTACACCTCCAGGCCGGGCATCTCCAGGCCG

IACACCAAAGTGACAGAAGTTCTTCCCTGGATTTACAGCAAGATGGAGAGCGAGGTGCGATTCAGAAAATCCTAA GCGGATCGTGGGGGGGGGCGTGGCCTCGGATAGCAAGTGGCCTTGGCAAGTGAGTCTGCACTTCGGCACCAC <u> CCACATCTGTGGAGGCACGCTCATTGACGCCCAGTGGGTGCTCACTGCCGCCCACTGCTTCTTCGTGACCCGG</u> GAGAAGGTCCTGGAGGGCTGGAAGGTGTACGCGGGCACCAGCAACCTGCACCAGTTGCCTGAGGCAGCTCC ATTGCCGAGATCATCATCAACAGCAATTACACCGATGAGGAGGACGACTATGACATCGCCCTCATGCGGCTGTC GCAGGTCAATCTCATCGACTTCAAGAAATGCAATGACTACTTGGTCTATGACAGTTACCTTACCCCAAGGATGAT AGACCTGCTGGATCACAGGCTTTGGCAAGACCAGGGAGACAGATGACAAGACATCCCCCTTCCTCCGGGAGGT ACCGAGGTTGCCCACAGGGATTTTGCCAACAGCTTCTCAATCTTGAGATACAACTCCACCATCCAGGAAAGCCT

>SGPR479 1 SEQID 46

GCTGCGCACTGCTTTCACAGGGACAAGAATATCAAAATCTATGACATGTACGTAGGCCTCGTAAACCTCAGGGTG CGGAGGTGACGTGGCCCTGGTGCAGCTGAAGACCCGCATTGTGTTTTCTGAGTCCGTGCTCCCGGTTTGCCTTG <u> GECCTETEGTCGECCCAGCATGGAGGGGAAAATCCTGGGCGGCGTCCCTGCGCCCGAGAGAGGGAAGTGGCCGTG</u> GCCGGCAACCACACCAGTGGTATGAGGTGAACAGGGTGATCCTGCACCCCCACATATGAGATGTACCACCCCAT CAACTCCAGAAGTGAACCTTACCAGTGCCAATTGCTGGGCTACGGGATGGGGACTAGTCTCAAAACAAGGTGAG ACCTCAGACGAGCTGCAGGAGGTGCAGCTCCCGCTGATCCTGGAGCCCTGGTGCCACCTGCTTACGGACACA TGTCCTACATCATGCCCGACATGCTGTGTGCTGGGGACATCCTGAATGCTAAGACCGTGTGTGAGGGGGGGACTCC SCAACCCTCTGTACCCTGGAGTGTATGCCAGTGTTTCCTATTTCTCAAAATGGATATGTGATAACATAGAAATCAC GCCCACTCCTGCTCAGCCAGCCCCTGCTCTCTCCAGCTCTGGGGCCCACTCTCAGCGTCTAATGGCCATGC GCAGGTCAGCGTGCACTACGCAGGCCTCCACGTCTGCGGCGGCTCCATCCTCAATGAGTACTGGGTGCTGTCA TGGCCCCTCCCCGGGTCGCAGCATTGGTCCACAGACAGCCAGAGAACCAGGGAATCTCCCTAACTGGCAGCGT GGGGGCCCACTTGTCTGTGAATTCAACCGCAGCTGGTTGCAGATTGGAATTGTGAGCTGGGGCCGAGGCTGCT 'GGCTGGCTGGTCAGTGCTGTGA

>SGPR489\_1\_SEQID\_47

CTGCAACTCTTTCGCTCCCCAAAGCTCCCAGTTGTGGGCAGAGTCTGGTTAAGGTACAGCCTTGGAATTATTTTA <u> ATGAĞTCTCAAAATGCTTATAAGCAGGAACAAGCTGATTTTACTACTAGGAATAGTCTTTTTTGAACGAGGTAAAT</u>

CTCACTATTGAAACTGTCATCATACATCCACATTTCTCCACCAAGAAACCAATGGACTATGATATTGCCCTTTTGAA ACATTITCAGTCGCATTCTTGGAGGAAGCCAAGTGGAGAAGGGTTCCTATCCCTGGCAGGTATCTCTGAAACAAA GGCAGAAGCATATTTGTGGAGGAAGCATCGTCTCACCACAGTGGGTGATCACGGCGGCTCACTGCATTGCAAAC GATGGCTGGAGCCTTCCAATTTGGCCACTTTGTGGGGCCCATATGTCTTCCAGAGCTGCGGGAGCAATTTGAGG STGTGGCACCTGGTGCAGTGAGCAGGATGTCATAGTCAGCGGGGCTGAGGGGGAAGCTGCACTTCCCAGAAAGC SCCGGAATAAGAAAGGGGCCTGGGACTCTGGCTGGTCAATTTGGGAGGCTCAGGTGGGAGGATCGCTTGAGTC CAGGGGCTTTCAGGCTATAGTCTCCTTCATTCCTAAAGCAGTATACCCCAGATTTAAACATCTCCATATCAGAGGAT CAGGAGTTCAAGACCAAGCCTAGGCAACAAGTGAGACTCTGTCTCACAAATAATTTCTTCAAAAAATTAGCCGG SGAAAATTTTGTGGAGAAAGCCTCCCTTCATCCATTCTTATTGGCTCTAATTCTCTAAGGCTGAAATTCGTCTCTG ATGCCACAGATTATGCAGCTGGGTTTAATCTTACCTATAAAGCTCTTAAACCAAACTACATTCCTGGTTGCAGTTA BATGTCCCCACCCTGTGCTGAGCCCCTCCAGCATCATGCTCATCAGCTTCCATTCAGATGAAAACGGGACCTG CTTAACTGTCCTTTTTGAAGAAGGTCTCATACAGAGTCTAAACTATCCTGAAAACTACAGTGACAAGGCTAACTGT CTGGTTTTATTTGTACAACTGCAGGCTGGGGCCGCTTAACTGAAGGTGGCGTCCTCTCACAAGTCTTGCAGGAA STGAATCTGCCTATTTTGACCTGGGAAGAGTGTGTGGCAGCTCTGTTAACACTAAAGAGGGCCCATCAGTGGGAA GACCTTTCTTTGCACAGGTTTTCCTGATGGAGGGAGAGGCGCATGTCAGGGAGATTCAGGAGGTTCACTCATGT CTCCACCTATATTATGAGAGCAAGCAACGGTGTGTCTGGACCCTGCTGGTACCAGAGGAAATGCATGTGTTGCT GAGTCAATGTTTCTGGAGACATGA

FIG. 1PP

>SGPR465 1 SEQID 48

ACTACACCATCCACGAGGACATGCTGTGCGCTGGGGACCTCATAACAGGAAAGGCCATTTGCCGAGTGAACTCC GATCATGGTGCACGCTGACTATAACGAGTTGCACCGCATGGGGAGTGACATCACCCTGCTGCAGCTGCACCATC AGGGGTCCCCTCGTCTGCCCATTAAATGGCACCTGGTTCCTGATGGGGCTGTCTAGTTGGAGCCTCGACTGCTG GATTACCGGATCCTGCTTGGGTATGACCAGCAAAGCCATCCCACAGAGCACAGCAGGAGGAGGAGATGACAGTGAATAA <u> GGGTGGCCTCTGCTGCTCCACTGCTTCCAAAGATGCATCTTCCCTCCACGGGCCCCCGCTGTCCACTAACCCATCT</u> CGGTGGCCATGGCAGGCCAGTCTCCTCTACCTAGGCGGGCACATCTGTGGAGCTGCCCTCATCGACAGCAAC1

CTCACCCGTCGGTCCCAGGGTCTTCACCAGGCTCCCCTACTTCACCAACTGGATCAGCCAGAAGAAGAGGGGAGA CATCGTCCACAAGCCCGGGCTCTGCGCAGCCCTTCTGGCTGCTCACATGTTCCTCCTGCTGCTGATTCTCCTGG GCACCCCTCCAGATCCCGCCTTGGCTCCTCCTCAGGAAACACCCCCCAGCCCTGGACAGCATGACCTCTCAGGG

>SGPR524 1 SEQID 49

TGGTCAAAGACATCACTGGCTTTGAAGGGAAAATTTCAAGCCCATATTACCCGAGCTACTATCCTCCAAAATGCA ACTTTTGGATTGTTTTTGTCATGCCACGTGCCAAAGGCCACATCTTCTGTGAAGACTGTGTTGCCGCCATCTTGAA AGTGTACCTGGAAATTTCAGACTTCTCTATCAACTCTTGGCATAGCACTGAAATTCTATAACTATTCAATAACCAAG STGTCCCTCAGGCCCAGCGTTGTGATGGAGTAAATGACTGCTTTGATGAAGTGATGAACTGTTTTGCGTGAGGC 4CAAAGGCTGCTCTCAGTACTTCTATGCAGAGCATCTGTCTCTCCACTACCCGCTGGAGATTTCTGCAGCCTCAG SCTCATATACGGAGGCTCTCAGGAATCCGGGCATATTTTGAGGTCATTCCAGAACAAAAGTGTGAAAACACAGTG CAGACAATITITCGAGTGCCCAGCCCTCTGGTTCACATTCAGCTCCAGTGCAGTTCAAGGCTTTCAGACAAGCCA CTTTTGGCAGAATATGGCAGTTACAACATCAGTCAACCCTGCCCTGTTGGATCTTTTAGATGCTCCTCCGGTTTAT <u> ATGGACAAAGAAAACAGCGATGTTTCAGCCGCACCTGCTGACCTGAAAATATCCAATATCTCAGTCCAAGTGGTC</u> AAAACAGAAAGCAAAGATGCTTTTTACTTTGCTGGGATGTTTCGCATCACCAACATTGAGTTTCTTCCCGAATACC CTGCCTTCTCCAAATITTATGAGCAGTCTGTTGTTGCAGATGTCAGCAGCAGCAACAAGAGGCGGCCTCCTTGTCC AGTGCCCAAAAGAAGCTGCCAGTGAGACGACCACCGTTGCCAGGGAGACGACTACCATTGCCAGGAAGACGAC CACCACAAAGACCCATTGGCAAAGCCAAACCCAAGAAGCAATCCAAGAAAAAAAGTTCCCTTTTGGAATGTACAAA <u> ATAAAATCATTCTCTTCACAGTATTTTATTCATCCTAGCAGTCATAGCCTGGACACTTCTGTGGCTGTATATCAG</u> SACAAAAGGAGTCCAGGGAATTTCTTTCAGTGTCACGGACTGTGCAGCAAGTGATAAACCTGGTTTATACAACAT SGACTCCATCCAGACAAGCATCATAAACCGGACCTCTGTGGGGGGGCTTGCAGGGACTGGCTGTGGACATGGAC STCAACCTGCCTGCAATACCAGCTCCTTCAGGCAGCATGGCCCTCTCATCTGTGATGGCTTCAGGGACTGTGAG CGAAGCCGACAACTGTGTCACTGACTCCCTGACCATTTACGACTCCCTTTTGCCCATCCGGAGCAGCATCTTGT GCTTTAGGAAACAAAATGCAAAATGTGATGGGACAGTGGATTGTCCAGATGGAAGTGATGAAGAAGGCTGCAC ACAGAATTTGTGAACCCACAAGAACATTAATGTCATTTGTTTCTACAAATAATCTCATGTTGGTGACATTTAAGTC1 4ATGGCCGGGATGAGCAAAACTGCACTCAAAGTATTCCATGCAACAACAGAACTTTTAAGTGTGGCAATGATATI

CATCACTTCTCGGATGCTCTGTGCAGGCATAATGTCAGGCAAGAGAGATGCCTGCAAAGGAGATTCGGGTGGAC CTTTATCTTGTCGAAGAAAAGTGATGGAAAATGGATTTTGACTGGCATTGTTAGCTGGGGACATGGATGTGGGC GAATGCCAAGTTTGTCTCCCCGGTGAGAAGTTGTGGTCCACGAGTACTATAACAGTCAGACTTTTGATTATGA TATTGCTTTGCTACAGCTCAGTATTGCCTGGCCTGAGACCCTGAAACAGCTCATTCAGCCAATATGCATTCCTCC AAAGGCTCCCTCGTTCTGCAGCAAGCGGAGGTAGAGCTCATTGATCAAACGCTCTGTGTTTCCACCTACGGGA1 3ACCAAACTTTCCTGGTGTTTACACAAGGGTGTCAAACTTTGTTCCCTGGATTCATAAATATGTCCCTTCTTTT

>SGPR422 SEQID 50

SGGACAAAGCAACACATATCAACTTAAGGACTTACGAGAGGACGACGGAAAATTTGGTGTATTCTTTGAAAATGTAC AAACCTGGAATGTTCTGTGCCGGATATATGGAAGGAATTTATGATGCCTGCAGGGGTGATTCTGGGGGGCCTTTA GTCACAAGGGATCTGAAAGATACGTGGTATCTCATTGGAATTGTAAGCTGGGGAGATAACTGTGGTCAAAAGGAC CTITCTITIGIGIGICACAGICCAGAGGAAGAIGGIGIGAAAGIAGAIGICATIAIGGIGITCCAGTICCCCTCTA AAGCAAGTTGTGGTAAACGAGTTGTTCCATTAAACGTCAACAGAATAGCATCTGGAGTCATTGCACCCAAGGCGG ATGACATTGAACAAAATTAAAGACCTTTTTGCAGGGAAAGGACAGTGGGATTTGGCACCCGAAGCAGAAATGCTG <u> AAGCCATGGATGATTGCCGTTCTCATTGTGTTGTCCCTGACAGTGGTGGCAGTGACCATAGGTCTCCTGGTTCAC</u> FGCATCCTTCCAACCAAATTTGACTGTCCACATCACAGGATTTGGAGCACTTTACTATGGTGGGGAATCCCAAAAT TTGTCACTGCAGCACACTGCTTCCAGAAGTATAAAAATCCACATCGATGGACTGTTAGTTTTGGAACAAAAATCAA CTGAACAAAGGGCAGTAAGAGAAGAAAATCCAAAGCATCTTAAATCAGAAGATAAGGAATTTAAGAGCCTTGC GGAGAGGCTCCAGGCCTGGGAGCAGGTCCTGCTGGTCACCAATGAGCTCATCAACAGGGGAGTTAACTGTCC CAATAAATGCCTCATCAGTTCAAGTTAATGTGGCCATGGTCAAGAATGGCAATGTGGGGCCAGGTTCCGGAGCA SCCTCCCTTAATGAAAAGAAATGTCAGAAGATTTATTATCCATGAGAAGTACCGCTCTGCAGCAAGAGAGTACGA TCCTAGTATTTGACCAAAAAAGGAGTACTATCATGGCTCCTTTAAAATTTTAGATCCACAAATCAATAACAATT **AAGCCTGGAGTCTACACACAAGTGACTTATTACCGAAACTGGATTGCTTCAAAAACAGGCATCTAA** 

1RR

>SGPR538 SEQID 51

GCAGTCTGTGGCTCCTGGGCGCTGGCCGTGGCCAGGCGTGGCCCTGGGCTTCCGGCACACGTGTGGGGG CCTGCTGCCTCTCAGCCCATTTCCGGGACCTTGCAGGATGAGGAGATAACTTTGAGCTGCTCAGAGGCCAGCGC STGCTGGAGCCTTGGGCATCTCAGACTCACTCACCACAAGGGAGTAAACCTCACTGACATCAAACTCAACAGTTC CTCTGTGCTAGCGCCACGCTGGGTGGTGACTGCTGCACATTGTATGCACAGTTTCAGGCTGGCCCGCCTGTCCA FGAGGAAGCTCTGCTCCCTGCACTCCCCAAAACAGTATCTTTCAGAATAAACAGCGAAGACTTCTTGCTGGAAGC <u> ACTTCTGGTCAAGTTGTTTCCCTCAGATGCTCTGAGTGTGGAGCGAGGCCCCTGGCTTCCCGGATAGTTGGTGG</u> GCTGGCGGGTTCATGCGGGGCTGGTCAGCCACAGTGCCGTCAGGCCCCCACCAAGGGGCTCTGGTGGAGAGGA CCAGGAGTTTGCTCAGCTCTCTCTAGACTGGGAGGCTTCCTGGAGGAGGCGTGGCAGCCCAGGAACAACTGC SCTACCTGGACGGAAGGGCTGATGCATGCCAGGGAGATAGCGGGGGCCCCCTAGTGTGCCCAGATGGGGACA TATCCCACACCCCCTCTACAGTGCCCAGAATCATGACTACGACGTCGCCCTCCTGAGGCTCCAGACCGCTCTC <u> GGGTGTCTGGCTGGGGCCACCCACCCTAGCCATACTTACAGCTCGGATATGCTCCAGGACACGGTGGTGCC</u> SCAAGTGAGGGATCAGCCACGCTGGCTCCTGGTCTGCCATGAGGGCTGGAGCCCCGCCCTGGGGCTGCAGAT GCAGTGCTGGGAGCCCTGGGGCTGCTGGCCGGTGCAGGTGTTGGCTCATGGCTCCTAGTGCTGTATCTGTGT <u> ATGAGCCTGATGCTGGATGACCAACCCCCTATGGAGGCCCAGTATGCAGAGGAGGGCCCAGGACCTGGGATCT</u> AACTTCTCAGACACTGTGGGCGCTGTGTGCCTGCCGGCCAAGGAACAGCATTTTCCGAAGGGCTCGCGGTGCT 

FIG. 1SS

>SGPR527\_1\_SEQID\_52

TCTCAGTCCTACCCAGGAAGAACCTGAAGATCTGGACTGCGGGCGCCCTGAGCCCTCGGCCCGCATCGTGGGG <u> ATGGCCCGGCACCTGCTCCTCCCCCTTGTGATGCTTGTCATCAGTCCCATCCCAGGAGCCTTCCAGGACTCAGC</u> GTGGCCGCCATCGTGGTGCCGGCCAACTACAGCCAAGTGGAGCTGGGCGCCCGACCTGGCCCTGCTGCGCCTG SECTCAAACECECAGCCGGGCACCTGGCCTTGGCAAGTGAGCCTGCACCATGGAGGTGGCCACATCTGCGGG SECACCECCTECTEGECCACCEGCTEGEGAGACGTCCAGGAGGCAGATCCTCTGCCTCTCCCCTGGGTGCTA SECTCCCTCATCECCCCTCCTGGGTCCTCTCCGCTGCTCACTGTTTCATGACGAATGGGACGCTGGAGCCCG 

SCTTTCTGGACCCGAACAGCTCCGACAGCCCACCCCGCGACCTCGACGCCTGGCGCGTGCTGCTGCTCGCC GGGTTCAGAGCCTGGGCCTGCCTTTCCCAGCCCCAGAAGACCCAGTCAGATCCCCAGGAGCCCAGGGAG SAGAACTGCACCATTGCCCTGCCTGAGTGCGGGAAGGCCCCGCGGCCAGGGGCCTGGCCCTGGGGGCCTGGGCCCAG TGTGGACGGAGAAACCGCCCTGGAGTTTTCACTGCTGTGGCTACCTATGAGGCATGGATACGGGAGCAGGTGAT SCGCTGCCCGGAGACCCGCCGCCGCGCGCTCTGCCTGCCTACCAGGAAAAGGAGGAGGTGGGCAGCTGCTGG 4CTCATGGCCCATGGATCAGCCATGTGACTCGGGGAGCCTACCTGGAGGACCAGCTAGCCTGGGATTGGGGCC CCTGCCCCCAGTGGCTCCCCACACTGGCCCACTGGAGGCAGCAATCTCTGCCCCCCAGAACTGGCCAAGGCC STGATGGGGAGGAGTGAGACACAGACTTGTCCCCCACACAGAGCATGGTGCCTGTGGCCTGCGGCTGG GCCCTCCCCATCAGCCCTGCCCATCTGTCTCCACCCGGGGGTATCCCCCCGGGGGCCAGCTGCTGGGTGTTG GGCTGGAAAGAACCCCAGGACCGAGTCCCTGTGGCTGCTGTCTCCCATCTTGACACAACGAATCTGTGACTG TCACTCTCCAGATATTGCCAGGGATGCTGTGTGCTGGCTACCCAGAGGGCCGCAGGGACACCTGCCAGGGTGA GTGATGGTGCCAGGATCCAGACCCTGCCATGGGGCGCTGGTGTCTGAAAGCTGGGTCTTGGCACCTGCCAGCT SCCGCGCGCGCGGGGGGGGCGCCCTGGTGCAGCACGAGACGCTTCGTGGGACAACGCCTCGGACCTGG SECTECTECAGCTGCGCACGCCCGTGAACCTGAGCGCGCGTTCGCGGCCCGTGTGCCTACCCCACCCGGAAC SECTECTCCAGTGGGGGTCCTGTGGCCTGGCTGGCAGAGGTGCATGTGGCTGGTGATCGAGTCTGCACTGG CAGCATCCGGCTGCCCCAGCACCTGGGACTCAGGCCCCCCCTGGCCCTCCTGGAGCTGAGCTCCCGGGTGGA SCTCTATCAGGGCATCCTGCCCCCTGGAACCCTCTGTGTCCTGTATGCAGAGGGGCCAGGAGAACAGGTGTGAG <u> ATGACCTCAGCACCGCCCTCCTGTGCCAGATGACGGAAGGGTCCTGGATCCTCGTGGGCATGGCTGTTCAAG</u> ACTACTTCCTGCCCGGGAGCCGCTGCCGCCTGGCCCGCTGGGCCGCGGGGAACCCGCGCGCTTGGCCCAGGC 36AGCCGGGAGCTGTTTGCTGCCATTGGTCCTGAAGAGGCCTGGATCTCCCAGACAGTGGGAGAGGCCAACT <u> 1 CGGGATCCCCGCATGCAGTCTACTTCCTGCTCCTGCTGACTCTCCTGATCCAGAGCTGA</u>

ATGCCATCCGTGCGCTTCGGGGGCCAACATCACTGCGGAGGCTTCCTGCTGCGAGCCCGCTGGGTGTCTCGG CCGCCCACTGCTTCAGCCACAGAGACCTCCGCACTGGCCTGGTGGTGGTGGGGCGCCCACGTCCTGAGTACTGC GCCATGGGGCTCGGGTTGAGGGGCTGGGGACGTCCTCTGCTGACTGTGGCCACCGCCCTGATGCTGCCGTG GGAGCCCACCCAGCAGGTGTTTGGCATCGATGCTCTCACCACACCCCCGACTACCACCCCATGACCCACGCC SGPR542 SEQID 53

AGGAGCTGCCGCCTGGACTGATGGAGGCCAAGGTCCGAGTGCTGGACCCGGACGTCTGCAACAGCTCCTGGAA GGGCCACCTGACACTTACCATGCTCTGCACCCGCAGTGGGGACAGCCACAGACGGGGGCTTCTGCTCGGCCGAC CCCAAGACCCCCGACGTGTACACGCAGGTGTCCGCCTTTGTGGCCTGGATCTGGGACGTGGTTCGGCGGGGGCA AACGACATCTGCCTGCTGCAGCTGAACGGCTCTGCTGTCCTGGGCCCTGCAGTGGGGCTGCTGAGGCTGCCAG TCCGGAGGGCCCCTGGTGTGCAGGAACCGGGCTCACGGCCTCGTTTCCTTCTCGGGCCTCTGGTGCGGCGAC GTCCCCAGCCCGGCCCCCTGGGACCACCAGGCCCCCAGGAGAAGCCGCCTGA

>SGPR551\_SEQID\_54

SGCCAGTGGACGATCCAGAACAGGAGGCTGTGTGGCTTGCGCATCCTGCAGCCCTACGCCGAGAGGATCCCCG CAAAGAGACAGCACATGCATCTCACTGCCCAAGGTCTGTGATGGGCAGCCTGATTGTCTCAACGGCAGCGATG <u> ATGCCCGTGGCCGAGGCCCCCCCAGGTGGCTGGCGGGCAGGGGGACGGAGGTGATGGCGAGGAAGCGGAGCCC</u> STECTECTGCCCTGCTCGTGCTGGCTTCGGCGGGGGTGCTACTCTGGTATTTCCTAGGGTACAAGGCGGAGG GTCCACAGTCAACAGCTCGGCTGCCGTCCCCTACAGGGCCGAGTACGAAGTGGACCCCGAGGGGCCTAGTGATC FGGACTACGGCTTGGCCCTCTGGTTTGATGCCTATGCACTGAGGAGGCAGAAGTATGATTTGCCGTGCACCCAG 4AGAGCAGTGCCAGGAAGGGGTGCCATGTGGGACATTCACCTTCCAGTGTGAGGACCGGAGCTGCGTGAAGAA IGATGGTCAGCCAGGTGTACTCAGGCAGTCTGCGTGTACTCAATCGCCACTTCTCCCAGGATCTTACCCGCCGG GGACCTCATGCTCAAACTCCGGCTGGAGTGGACGCTGGCAGAGTGCCGGGACCGACTGGCCATGTGACGTG CCTCAGCACCCCGTACTTCCCCAGCTACTACTCGCCCCAAACCCACTGCTCCTGGCACCTCACGGTGCCTCTC GETGGCCACGGCCGGGATCACCATCAACTTCACCTCCCAGATCTCCCTCACCGGGCCCGGTGTGCGGGTGCC GAATCTAGTGCCTTCCGCAGTGAAACCGCCAAAGCCCAGAAGATGCTCAAGGAGCTCATCACCAGCACCCGCCT <u> COGTECAECCEGTEGTCTTCCAGGCCTGTGAAGTGAACCTGACGCTGGACAACAGGCTCGACTCCCAGGGCGT</u> STGGCGTCGGGGGCCATCATGGCGGTCGTCTGGAAGAAGGGCCTGCACAGCTACTACGACCCCTTCGTGCTCT CTGGAAGCCAGTGTGAAAGACATAGCTGCATTGAATTCCACGCTGGGTTGTTACCGCTACAGCTACGTGGGCCA SGGCCAGGTCCTCCGGCTGAAGGGGCCTGACCACCTGGCCTCCAGCTGCCTGTGGCACCTGCAGGGCCCCAA CTATGGCTTGTACAACCAGTCGGACCCCTGCCCTGGAGAGTTCCTCTGTTCTGTGAATGGACTCTGTGTCCTG SCTGTGATGGGGTCAAGGACTGCCCCAACGGCCTGGATGAGAGAACTGCGTTTGCAGAGCCACATTCCAGTG AGAGGGGATGTTCAAGGCCTGTGAGGACTCCAAGAGAAAAGCCCGGGGGTACCTCCGCCTGGTGCCCCTGTTT SCAAATCCCCGAGCACCGCCGGCTGATGCTGAGCCCCGAGGTGGTGCAGGCACTGCTGGTGGAGGAGCTGCT SCCGGGCCCCTGGAGAGAGAGGCTCATCACCTCGGTGTACGGCTGCAGCCGCCAGGAGCCCGTGGTGGAGGTT SCCCAACCCGCAGTGTGATGGGCGGCCCGACTGCAGGGACGGCTCGGATGAGGAGCACTGTGACTGTGGCCT

FIG. 1UU

CCAGGGCCCCTCCAGCCGCATTGTTGGTGGAGCTGTGTCCTCCGAGGGTGAGTGGCCATGGCAGGCCAGCCTC rccaggaggacagcatggcctccacggtgctgtggaccgtgttcctgggcaaggtgtggcagaactcgcgctg SACGTGGCGCTGCTGCAGCTCGACCACCCGGTGGTGCGCTCGGCCGCCGTGCGCCCCGTTTGCCTGCCCGCG **ATCAGCAACGCTCTGCAGAAAGTGGATGTGCAGTTGATCCCACAGGACCTGTGCAGCGAGGCCTATCGCTACCA** CCGCTGGTGTGCAAGGCACTCAGTGGCCGCTGGTTCCTGGCGGGGCTGGTCAGCTGGGGGCCTGGGGCTTGTGGC CAGGTTCGGGGTCGACACATCTGTGGGGGGGCCCTCATCGCTGACCGCTGGGTGATAACAGCTGCCCACTGCT GCCTGGAGAGGTGTCCTTCAAGGTGAGCCGCCTGCTCCTGCACCGTACCACGAAGAGAGACAGCCATGACTAC CGGCCTAACTACTTCGGCGTCTACACCCGCATCACAGGTGTGATCAGCTGGATCCAGCAAGTGGTGACCTGA

SGPR451 SEQID 55

A GTTA C G G A C A T G G C T G T G G A G G G T T T C C T G G T C T A T T G G G C C A T C T C T A C C A A A G T G G C T G CAGCCTATTTGCCTACCTTTTGATGTTTTCCAAATCCTGGACGGAAACACAAAGTGTTTTATAAGTGGCTGGGGAA SAACAAAAGAAGAAGGTAACGCTACAAATATTTTACAAGATGCAGAAGTGCATTATATTTCTCGAGAGATGTGTAA FGATTGGAACTAATAATATACATGGACGCTATCCTCATACCAAGAAGATAAAAATTAAAGCAATCATTATTCATCCA TCTGAGAGGAGTTATGGGGGAATAATTCCTAACACTTCATTTTGTGCAGGTGATGAAGATGGAGCTTTTGATACT \GGATTGTGGAACAGCACCGCTTAAGGATGTGTTGCAAGGGTCTCGGATTATAGGGGGCACCGAAGCACAAGCT **AACTTCATTTTGGAATCTTATGTAAATGATATTGCACTTTTTCACTTAAAAAAAGCAGTGAGGTATAATGACTATAT** GGAGCTCTCACTTATACTCAGACCACTACTCGCCCTCTGGAAGGCACAGGCTCGGCCCCTCGCCGGAACCGGC GCAGGGGTGACAGTGGGGGACCATTAATGTGCTACTTACCAGAATATAAAAGATTTTTTGTAATGGGAATTACC <u> GACCTGCCGCCATCTTGCTCACCAGCCTCCAAAATGCGGCTGGGGCTCCTGAGCGTGGCGCGTGTTGTTGTGG</u> **TATGTTTTGTCATCTTACTAGCAACAACATAA** 

**1VV** 

>SGPR452\_1\_SEQID\_56

TGGGCAGCAGCTTGAAGTGGGCGTGGGCCAAGCCCTCTGGGATGCCCGTCCCAGAGAATGACCTGGTGGGCAT AGCCCCCGCAGCCCCAGGACCCCTGACTGTAGGCTCCAGGCCTCCCTGGAAGCCCTGGCCACGCTCGCCCCG CAGCCCTCAGACTGGCTGTGCTTCGCGGATCTTGGCTGGTTCGAGGCTGATGGAGCTGCCCACTCCATGGGCC

SGTGCTGGAGAACGCCGTCTGTGAGCAGCCCTACCGCAACGCCTCAGGGCACACTGGCGACCGGCAGCTCATC TCTGGAAGGACACCGACCCGTCCATCTACCGGATCCACGCTGGGGACGTGTATCTCTACGGGGGCCGGGGG SCAGGCTGCGGGGGTCCTGGCGCCTGGTGGGGGTGGTCAGCTGGGGGCTACGGCTGTACCCTGCGGGACTTTC CTGGATGACATGCTGTGTGCCGGCAGCGAGGGCCGAGACTCCTGCTACGGTGACTCCGGCGGCGCCCTCTGGTCT TGTGGGGGGCCACATGCCCCCCCCGGGGAAGTGGCCGTGGCAGGTCAGCCTGAGGGTTACAGCTACACTG FGCTGAACGTCAGCCGGATCATCGTCCACCCCAACTATGTCACTGCGGGGCTGGGGTGCGGATGTGGCCTTGCT SCAGCTGCCGGGGTCACCTCTCCCCAGAGTCGCTGCCGCCGCCTACCGCCTGCAGCAGGCGAGTGTGCA GGCCTCCTGGGCGCACATCTGTGGGGGCTCCCTCATCCACCCCCAGTGGGTGCTGACTGCTGCCGCCACTGCATT SCGGCGTCTACACCCACGTCCAGATCTACGTGCTCTGGATCCTGCAGCAAGTCGGGGAGTTGCCCTGA

OSSESIS CEECI

SGPR504 SEQID 57

<u> ATCGGGGGCCACGAGGTGACCCCCCCACTCCAGGCCCTACATGGCATCCGTGCGCTTCGGGGGGCCAACATCACT</u> SCGGAGGCTTCCTGCTGCGAGCCCGCTGGGTGGTCTCGGCCGCCCAGTGCTTCAGCCACAGG

SGPR469 SEQID 58

GGAGATTĈTGGGĞGGCCCCTGGTCTGTGAATTAAATGGCACATGGGTCCAGGTGGGGATTGTGAGCTGGGGCA 

SGPR400 SEQID 59

SCCCAGTTTCTGTTAGCCCTGTTGTCCTGCTCATCTGCCTTCCCTCATCTGAAGTCTACCTGAAGAAGAATACAAC <u> 2AGCATCCGCCAGGGCTTGATTCACGTCTGCTCAGATACCCTCATCTCAGAGGAGTGGGTGCTGACAGTGGCGA</u> STGGGCGGCCCACAGTCTCATCTGGTATTGCCTCAGGCTTGGGGGCTAGTGTGGGGCCAGTGGCCCTGGCAGG CACGGGCACTGCTGGCTATGCATTCACGCTGCTCCTTCTGCTGGGGATTTCGGGTGAGCCCCCCAGAATGGGTC1 SAAAGTGCCCCTCATTGATCTCCAGACATGCGGTGACCACTATCAAAATGAAATCTTGCTGCACGGAGTTGAGCT STCCTGCTGGGTGACTGGATGGGGCTATACTGGAATATTCCAATATATCAAGCGTTCTTATACACTGAAGGAGCT ICTGCTTCCCATTATCCCCCCCACCCTGATTTCCAAGCAAACACATCTAGTGCCATCGCTGTGGTAGAACTGCCCT SAGAGTCAGGTTCTTCCCGATTTGACAGCAGGGGACGCCGCAGACCCCCCCAATTCCTCCCTTGGGTCCTGGAG CATCATCAGTGAAGCTATGATCTGCTCCAAGCTCCCAGTGGGCAGATGGATCAGTGTACTGTAAGAATCCACC <u> ATTICICCITICITATAGAGCATCACTCCTTGCTGTTGTAACACACAGATCCAATAATAGTCGTGGĠCGAGCTTTI</u> SCTCAGGCACCTTTCACAGGCCTTGCCTTCCCCAGTGA DJEBSGIS "CSEGI

>SGPr397 SEQID 60

GLIHMFSIGRSYEGRCLFILKLGRRSRLKRAVWIDCGIHAREWIGPAFCQWFVKEALLTYKSDPAMRKMLNHLYFYIMP MKCLGKRRGQAAAFLPLCWLFLKILQPGHSHLYNNRYAGDKVIRFIPKTEEEAYALKKISYQLKVDLWQPSSISYVSEG VFNVDGYHFSWTNDRFWRKTRSRNSRFRCRGVDANRNWKVKWCGKFGTNWDPDPKVSAGFTLQNMSPEDSHGR TVTDVHIPQNGSRALLAFLQEANIQYKVLIEDLQKTLEKGSSLHTQRNRRSLSGYNYEVYHSLEEIQNWMHHLNKTHS

>SGPr413 SEQID 61

WTTDRLWRKSRSPHNNGTCFGTDLNRNFNASWCSIGASRNCQDQTFCGTGPVSEPETKAVASFIESKKDDILCFLTM MKPLLETLYLLGMLVPGGLGYDRSLAQHRQEIVDKSVSPWSLETYSYNIYHPMGEIYEWMREISEKYKEVVTQHFLGV HSYGQLILTPYGYTKNKSSNHPEMIQVGQKAANALKAKYGTNYRVGSSADILYASSGSSRDWARDIGIPFSYTFELRD TYETHPMYYLKISQPSGNPKKIIWMDCGIHAREWIAPAFCQWFVKEILQNHKDNSSIRKLLRNLDFYVLPVLNIDGYIYT SGTYGFVLPEAQIQPTCEETMEAVLSVLDDVYAKHWHSDSAGRVTSATMLLGLLVSCMSLL

>SGPr404 SEQID 62

YPHESQLPEEWENNRESLIVFMEQVHRGIKGLVRDSHGKGIPNAIISVEGINHDIRTANDGDYWRLLNPGEYVVTAKAE RNEMTTTDDLDFKHHNYKEMRQLMKVVNEMCPNITRIYNIGKSHQGLKLYAVEISDHPGEHEVGEPEFHYIAGAHGNE LLWEAEDRQNVPRKVPNHYIAIPEWFLSENATVAAETRAVIAWMEKIPFVLGGNLQGGELVVAYPYDLVRSPWKTQEH VLGRELLLLLVQFVCQEYLARNARIVHLVEETRIHVLPSLNPDGYEKAYEGGSELGGWSLGRWTHDGIDINNNFPDLN1 MVSNDSHTWVTVKNGSGDMIFEGNSEKEIPVLNELPVPMVARYIRINPQSWFDNGSICMRMEILGCPLPDPNNYYHR PTPDDHVFRWLAYSYASTHRLMTDARRRVCHTEDFQKEEGTVNGASWHTVAGSLNDFSYLHTNCFELSIYVGCDK **GFTASTKNCMVGYDMGATRCDFTLSKTNMARIREIMEKFGKQPVSLPARRLKLRGRKRRQRG** 

>SGPr536 1 SEQID 63

GSWAFSTAAVASDRVSIHSLGHMTPVLSPQNLLSCDTHQQGGCRGGRLDGAWWFLRRRGVVSDHCYPFSGRERDE CNRTVSDCCPDFWDFCLGVPPFFPPIQGCMHGGRIYPVLGTYWDNCNRCTCQENRQWQCDQEPCLVDPDMIKAIN AGPAPPCMMHSRAMGRGKRQATAHCPNSYVNNNDIYQVTPVYRLGSNDKEIMKELMENGPVQALMEVHEDFFLYK MWRCPLGLILLLPLÄGHLALGAQQGRGRRELAPGLHLRGIRDAGGRYCQEQDLCCRGRADDCALPYLGAICYCDLF QGNYGWQAGNHSAFWGMTLDEGIRYRLGTIRPSSSVMNMHEIYTVLNPGEVLPTAFEASEKWPNLIHEPLDQGNCA **3GIYSHTPVSLGRPERYRRHGTHSVKITGWGEETLPDGRTLKYWTAANSWGPAWGERGHFRIVRGVNECDIESFVL** 

KYFMSPTLTMRLAGLSQITNQLHTFNDVCNNESLVSDTETSIAKELADWLISNNVVEHIFGPNLHIEIIKQCQVILNFLAAE MLMAFQNISDEQSFKAQSDHRSRHEVSHYSMWLLVSWAHCCSLVKSSLADSDHLQDWLKKLTLLIPETAVRHESCSG EYDLIGVTVHTGTADGGHYYSFIRDIVNPHAYKNNKWYLFNDAEVKPFDSAQLASECFGGEMTTKTYDSVTDKFMDFS ATHIAQGSQESCITRTGDFLGETIGNELFNCRQFIGPQHHHHHHHHHHHHDGHMVDDMLSADDVSCSSSQVSAKSEK /IQRLRPVHAHLYLQPGMEDGSDDMDTSVEDIGGRSCVTRFVRTLLLIMEHGVKPHSKHLTEYFAFLYEFAKMGEEES **HSSRAAAYDLLVEMVKGSVENYRLIHNWVMAQHMQSHAPYKWDYWPHEDVRAECRFVGLTNLGATCYLASTIQQLY** SKQSSFASLLNTNIPIGNKKEEEELRRTAPSPWSPAASPQSSDNSDTHQSGGSDIEMDEQLINRTKHVQQRLSDTEE \_CVAKIFQIQFPLYTAYKHNTHPTIEDISTQESNILGAFCDMNDVEVPLHLLRYVCLFCGKNGLSLMKDCFEYGTPETLP **3RLSTQHIDCIWAAAQLKHCSRYIHDLFPSLIKNLDPVPLRHLLNLVSALEPSVHTEQTLYLASMLIKALWNNALAAKAQ** SMQGSSDETANSGEDGSSGPGSSSGHSDGSSNEVNSSHASQSAGSPGSEVQSEDIADIEALKEEDEDDDHGHNPP DIVQDEDAVNLSEGLINEAEKLLCSLVCWFTDRQIRMRFIEGCLENLGNNRSVVISLRLLPKLFGTFQQFGSSYDTHWI1 DALARHLADCIRSREILDHQDGNVEDDGLTGLLRLATSVVKHKPPFKFSREGQEFLRDIFNLLFLLPSLKDRQQPKCKS EMSPELKNTVKSLFGGVITNNVVSLDCEHVSQTAEEFYTVRCQVADMKNIYESLDEVTIKDTLEGDNMYTCSQCGKKV RAEKRACFKKLPRIXSFNTMRYTFNMVTMMKEKVNTHFSFPLRLDMTPYTEDFLMGKSERKEGFKEVSDHSKDSESY FEKTHSAYMLFYKRMEPEEENGREYKFDVSSELLEWIWHDNMQFLQDKNIFEHTYFGFMWQLCSCIPSTLPDPKAVS KSSCGTDLRNRKLESQAGICLGDSQGTSERNGTSSGTGKDLVFNTESLPSVDNRMRMLDACSHSEDPEHDISGEMN NQQAQLQEFGQSNRKGEFPGGLMGPVRMISSGHELTTDYDEKALHELGFKDMQMVFVSLGAPRRERKGEGVQLPA **QFLLSLQAISTMVHFYMGTKGPENPQVEVLSEEEGGEEEEEEDILSLAEEKYRPAALEKMIALVALLVEQSRSERHLTL** SQTDMAALTGGKGFPFLFQHIRDGINIRQTCNLIFSLCRYNNRLAEHIVSMLFTSIAKLTPEAANPFFKLLTMLMEFAGG MWAEKELNMMKLFFDNLVYYIQTVREGRQKHALYSHSAEVQVRLQFLTCVFSTLGSPDHFRLSLEQVDILWHCLVED SECYDDALHWFLNQVRSKDQHAMGMETYKHLFLEKMPQLKPETISMTGLNLFQHLCNLARLATSAYDGCSNSELCG MDQFWGIALRAQSGDVSRAAIQYINSYYINGKTGLEKEQEFISKCMESLMIASSSLEQESHSSLMVIERGLLMLKTHLE LYKLSLSGLDGGDSINRSFLLLAASTLLKFLPDAQALKPIRIDDYEEEPILKPGCKEYFWLLCKLVDNIHIKDASQTTLLDL MIPEARQAVFTAKYSEDMKHKTTLLELQKMFTYLMESECKAYNPRPFCKTYTMDKQPLNTGEQKDMTEFFTDLITKIE -MTAKLSTSFVLETFIHSKEKPTMLQWIELLTKQFNNSQAACEWFLDRMADDDWWPMQILIKCPNQIVRQMFQRLCIH FLIAHAFITVVSNIRIWLHIPAVMQHIIPFRTYVIRYLCKLSDQELRQSAARNMADLMWSTVKEPLDTTLCFDKESLDLAF NMADFDGEESGCEEELVQINSHAELTSHLQQHLPNLASIYHEHLSQGPVVHKHQFNSNAVTDINLDNVCKKGNTLLW MCENCADLVEVLNEISDVEGGDGLQLRKEHTLKIFTYINSWTQRQCLCCFKEYKHLEIFNQVVCALINLVIAQVQVLRD AFRRFAYHLRQWQIEGTGISSHLKALSDKQSLPLRVVCQPAGLPDKMTIEMYPSDQVADLRAEVTHWYENLQKEQI SCLPPPQKDNIPMLLLLQEPHLTTLFDLLEMLASFKPPSGKVAVDDSESLRCEELHLHAENLSRRVWELLMLLPTCPN >SGPr414 SEQID 64

FIG. 2C

LORKDVKOALIOWOERIEFAHKLLTLLNSYSPPELRNACIDVLKELVLLSPHDFLHTLVPFLOHNHCTYHHSNIPMSLGP YFPCRENIKLIGGKSNIRPPRPELNMCLLPTMVETSKGKDDVYDRMLLDYFFSYHQFIHLLCRVAINCEKFTETLVKLSV LVAYEGLPLHLALFPKLWTELCOTOSAMSKNCIKLLCEDPVFAEYIKCILMDERTFLNNNIVYTFMTHFLLKVQSQVFSE LQEQEAKERKTKDDEGATPIKRRVSSDEEHTVDSCISDMKTETREVLTPTSTSDNETRDSSIIDPGTEQDLPSPENSS PPGMPPFASYILQRIWEVIEYNPSQCLDWLAVQTPRNKLAHSWVLQNMENWVERFLLAHNYPRVRTSAAYLLVSLIPS TYFMDLWNLFQPKLSEPAIATNHNKQALLSFWYNVCADCPENIRLIVQNPVVTKNIAFNYILADHDDQDVVLFNRGMLP ANCANLISTLITNLISQYQNLQSDFSNRVEISKASASLNGDLRALALLLSVHTPKQLNPALIPTLQELLSKCRTCLQQRNS **AYYGILRLCCEQSPAFTRQLASHQNIQWAFKNLTPHASQYPGAVEELFNLMQLFIAQRPDMREEELEDIKQFKKTTISC** VKEYRMEVPSSFSEDMSNIRSQHAEEQSNNGRYDDCKEFKDLHCSKDSTLAEEESEFPSTSISAVLSDLADLRSCDG NSFRQMFRSTRSLHIPTRDLPLSPDTTVVLHQVYNVLLGLLSRAKLYVDAAVHGTTKLVPYFSFMTYCLISKTEKLMFS YLRCLDGRSCWTTLISAFRILLESDEDRLLVVFNRGLILMTESFNTLHMMYHEATACHVTGDLVELLSIFLSVLKSTRPY QALPSQDPEVALSLSCGHSRGLFSHMQQHDILDTLCRTIESTIHVVTRISGKGNQAAS

COBSSELT . OSESOL

SGPr430\_SEQID\_65

KFNRLPRVLILHLKRYSFNVALSLNNKIGQQVIIPRYLTLSSHCTENTKPPFTLGWSAHMAMSRPLKASQMVNSCITSPS FLSIDKVPSKDAEEMRLFLDAVHQNRLPAAMKPSQGSGSFGAILGSRTSQKETSRQLSYSDNQASAKRGSLETKDDIP FRKVLGNPGRGSIKTVAGSGIARTIPSLTSTSTPLRSGLLENRTEKRKRMISTGSELNEDYPKENDSSSNNKAMTDPSR DVFDMEYTEAEAEELKRNAETGNLPHSYRLISVVSHIGSTSSSGHYISDVYDIKKQAWFTYNDLEVSKIQEAAVQSDRD MSPLKIHGPIRIRSMQTGITKWKEGSFEIVEKENKVSLVVHYNTGGIPRIFQLSHNIKNVVLRPSGAKQSRLMLTLQDNS LVKKDICNSETKKDLLKKVKNAISATAERFSGYMQNDAHEFLSQCLDQLKEDMEKLNKTWKTEPVSGEENSPDISATR AYTCPVITNLEFEVQHSIICKACGEIIPKREQFNDLSIDLPRRKKPLPPRSIQDSLDLFFRAEELEYSCEKCGGKCALVRH KYLTSSREKQLSLKQSEENRTSGGLLPLQSSSFYGSRAGSKEHSSGGTNLDRTNVSSQTPSAKRSLGFLPQPVPLSV KKLRCNQDYTGWNKPRVPLSSHQQQQLQGFSNLGNTCYMNAILQSLFSLQSFANDLLKQGIPWKKIPLNALIRRFAHL TPSKKFTFKSKSSLALCLDSDSEDELKRSVALSQRLCEMLGNEQQQEDLEKDSKLCPIEPDKSELENSGFDRMSEEEL -AAVLEISKRDASPSLSHEDDDKPTSSPDTGFAEDDIQEMPENPDTMETEKPKTITELDPASFTEITKDCDENKENKTP EGSQGEVDWLQQYDMEREREEQELQQALAQSLQEQEAWEQKEDDDLKRATELSLQEFNNSFVDALGSDEDSGNE RSGYIFFYMHKEIFDELLETEKNSQSLSTEVGKTTRQAS

>SGPr496\_1\_SEQID\_66

FNNDGVCCCLQKRGPVNITSVCVSPRTLQISVFVLSEKYEGIVKFESDELPFGVIGSNIGDAHFQEFRAGISWKPVVDP DDPIPQFPDCCSSSSSRIPSVSVLVAVPLVAGHKGQAFIERMLGCFKELKQELTQEGPGGGHPRSAWPPRRHAQWP MTLLAPWYTGPMIPMDVNEPSSVTTAPTLSSSLQHISSFLATGKKLSLHFGHPRECEVTRIDDKNRRGLEDSEPGAKL

PGRRAELKLEPEPEPVREAEQEPKQELEDENPARSGGGGNSDEVPPTLPSDPPRPPDPSPRRSRAPRRPRPRPQ LQASTSTEVSHQQCSVPGLGEKFPTWETTKPELELLGHNPRRRRITSSFTIGLRGLINLGNTCFMNCIVQALTHTPILRD FFLSDRHRCEMPSPELCLVCEMSSLFRELYSGNPSPHVPYKLLHLVWIHARHLAGYRQQDAHEFLIAALDVLHRHCKG **ASSKESRMNGQLQLPTNSGNNENKYSLFAVVNHQGTLESGHYTSFIRHHKDQWFKCDDAVITKASIKDVLDSEGYLLF** KSCICHVCGTHLNRLHSCLSCVFFGCFTEKHIHEHAETKQHNLAVDLYYGGIYCFMCKDYVYDKDIEQIAKEEQGEALK TLTDCLRRFTRPEHLGSSAKIKCGSCQSYQESTKQLTMNKLPVVACFHFKRFEHSAKQRRKITTYISFPLELDMTPFM TRLRTPPQPRPPPPPPPRPRRGPGGCLDVDFAVGPPGCSHVNSFKVGENWRQELRVIYQCFVWCGTPETRKSKA DDVGKAANNPNHCNCIIDQIFTGGLQSDVTCQACHGVSTTIDPCWDISLDLPGSCTSFWPMSPGRESSVNGESHIPGI PEPCEQGEEPPPVEAEEVEEAETAEKAERKVEAEAKVEGKAEAAGKAEAAGKVDATEKVETAGKVDAAGKVETAEG YHKQVLEHESEKVKEMNTQAY

DOSSOLS OSSOL

>SGPr495 SEQID 67

-NISNNLCFLEGKHLRSYSPQNAFQTLSQSYITTSKECSIQSCLYQFTSMELLMGNNKLLCENCTKNKQKYQEETSFAE MRVKDPTKALPEKAKRSKRPTVPHDEDSSDDIAVGLTCQHVSHAISVNHVKRAIAENLWSVCSECLEERRFYDGQLVL DSEPSESESASKQTGLFRSSSGSGVQPDGPLYPLSAGKLLYTKETDSGDKEMAEAISELRLSSTVTGDQDFDRENQP ASILKAFNNPTTKTADDETRKKVKISTVKDPFIDISLPIIEERVSKPLLWGRMNKYRSLRETDHDRYSGNVTIENIHQPRA KKVEGVYTNARKQLLISAVPAVLILHLKRFHQAGLSLRKVNRHVDFPLMLDLAPFCSATCKNASVGDKVLYGLYGIVEH **AKKHSSSKDKSQLIHDRKCIRKLSSGETVTYQKNENLEMNGDSLMFASLMNSESRLNESPTDDSEKEASHSESNVDA QTSAFSRIMKLCEEKCETDEIQKGGKCRNLSVRGITNLGNTCFFNAVMQNLAQTYTLTDLMNEIKESSTKLKIFPSSDS** QLDPLVVELSRPGPLTSALFLFLHSMKETEKGPLSPKVLFNQLCQKAPRFKDFQQQDSQELLHYLLDAVRTEETKRIQ SGSMREGHYTAYVKVRTPSRKLSEHNTKKKNVPGLKAADSESAGQWVHVSDTYLQVVPESRALSAQAYLLFYERVL TSDIWLCLKCGFQGCGKNSESQHSLKHFKSSRTEPHCIIINLSTWIIWCYECDEKLSTHCNKKVLAQIVDFLQKHASKT

>SGPr407 SEQID 68

WEYPVPYFRSPNRTLIPERIWSNPLLVLVIAYKTVSWPRQQLLAKQANKWMPFVIPSKTLPWDPLELKICYQQNRPYPS PDPSNFPTFLRCLNAFSAAVFYLPQPSWHKPEGLKPAGYPRVPDIPYGSGYTLKSTTEAAGLHQSLPMVQLPLHPTKG KAGKLFQPDAHGFLVKKVHAPTRGIVFIMEPRQLGGKGSLSKLQPACALGGMNSGMEPQKSAPFAAGKGLAPPLPVC VSLRDCLSLFTKEEELELENASGTLPVTKSEVLSTSCVPFGTTQAASTVATTQPCASARLVGTFTMTLVSPLNTLRDTE GIELTVMKALVLDILFKASTDIILFNHDSSSGNKWRKLPEPGGLEKKHEELRLRPLKEEYHWLVLVPLKLTGSPHRWRP RKRALASCSWCLQRVTMRRVMGVQDKAGNRNQMLLLGQRPVIGDTVSNSQTTRDKACRRPPSHSVFTQSSFWACL SALLKESELNDADWANLMWKRYLEEQEDSKMVDLFVGQMKSYLKCQACGYHSMTFKVFFFCDLSLTIPKKGFAGGK DPDLFFYGHQSYWMKAHLNDLILREGPVTQMAQSFYWGFPAGGNLSALEMLPDGPAPRTFLQKKSCLFPLFSYILLH

NLRFKLRVYKFEEELWSRAGLGKKSDNHSSRQMPWGAAGVACQHPCKLPRIVAELTPPKLSFGFLNTVQSSVLPTSL SOFFLNDSQPEEAIPPQSLLPGSPRTNSFPKDKFVPKDKLKVILSLLTMYELDRLF

>SGPr453 SEQID\_69

NIFHGQLLSQVTCLACDNKSNTIEPFWDLSLEFPERYQCSGKDJASQPCLVTEMLAKFTETEALEGKIYVCDQCNSKRR **RRILMGKIFRTWFEQSPIGRKKQEEPFQEKIVVKREVKKRRQELEYQVKAELESMPPRKSLRLQGLAQSTIIEIVSVQVP** VMHHGKGFGSGHYTAYCYNSEGGFWVHCNDSKLSMCTMDEVCKAQAYILFYTQRVTENGHSKLLPPELLLGSQHPN RFSSKPVVLTEAQKQLMICHLPQVLRLHLKRFRWSGRNNREKIGVHVGFEEILNMEPYCCRETLKSLRPECFIYDLSAV VFCYLCDDYVLNDNATGDLKLLRRTLSAIKSQNYHCTTRSGRFLRSMGTGDDSYFLHDGAQSLLQSEDQLYTALWHR MTASEKTRSCKHPPVTDTVVYQMNECQEKDTGFVCSRQSSLSSGLSGGASKGRKMELIQPKEPTSQYISLCHELHTL MLAMDTCKHVGQLQLAQDHSSLNPQKWHCVDCNTTESIWACLSCSHVACGRYIEEHALKHFQESSHPVALEVNEMY FQVMWSGKWALVSPFAMLHSVWRLIPAFRGYAQQDAQEFLCELLDKIQRELETTGTSLPALIPTSQRKLIKQVLNVVN AQTPASPAKDKVLSTSENEISQKVSDSSVKRRPIVTPGVTGLRNLGNTCYMNSVLQVLSHLLIFRQCFLKLDLNQWLA

>SGPr445\_SEQID\_70

MRVKDPTKALPEKAKRSKRPTVPHDEDSSDDIAVGLTCQHVSHAISVNHVKRAIAENLWSVCSECLEERRFYDGQLVI QTSAFSRIMKLCEEKCETDEIQKGGKCRNLSVRGITNLGNTCFFNAVMQNLAQTYTLTDLMNEIKESSTKLKIFPSSDS TSDIWLCLKCGFQGCGKNSESQHSLKHFKSSRTEPHCIIINLSTWIIWCYECDEKLSTHCNKKVLAQIVDFLQKHASKT **QLDPLVVELSRPGPLTSALFLFLHSMKETEKGPLSPKVLFNQLCQKRVHLHL** 

FIG. 2E

>SGPr401 1\_SEQID\_71

MTVRNIASICNMGTNASALEKDIGPEQFPINEHYFGLVNFGNTCYCNSVLQALYFCRPFRENVLAYKAQQKKKENLLTC EAQKRMRVKKLPMVLALHLKRFKYMEQLRRYTKLSYRVVFPLELRLFNTSSDAVNLDRMYDLVAVVVHCGSGPNRGH LADLFHSIATQKKKVGVIPPKKFISRLRKENDLFDNYMQQDAHEFLNYLLNTIADILQEEKKQEKQNGKLKNGNMNEPA ENNKPELTWVHEIFQGTLTNETRCLNCETVSSKDEDFLDLSVDVEQNTSITHCLRDFSNTETLCSEQKYYCETCCSKQ YITIVKSHGFWLLFDDDIVEKIDAQAIEEFYGLTSDISKNSESGYILFYQSRE

>SGPr408 SEQID 72

SDYVSQSYSYSSILNKSETGYVGLVNQAMTCYLNSLLQTLFMTPEFRNALYKWEFEESEEDPVTSIPYQLQRLFVLLQT <u>MVPGEENQLVPKEAPLDHTSDKSLLDANFEPGKKNFLHLTDKDGEQPQILLEDSSAGEDSVHDRFIGPLPREGSVGS1</u> SKKRAIETTDVTRSFGWDSSEAWQQHDVQELCRVMFDALEQKWKQTEQADLINELYQGKLKDYVRCLECGYEGWRI

GLEKNSLIYELFSVMAHSGSAAGGHYYACIKSFSDEQWYSFDDQHVSRITQEDIKKTHGGSSGSRGYYSSAFASSTNA VNNDRSTSSVDSDILSSSHSSDTLCNADNAQIPLANGLDSHSITSSRRTKANEGKKETWDTAEEDSGTDSEYDESGKS KTWKNPGTVFLDYHIYEEDINISSNWEVFLEVLDGVEKMKSMSQLAVLSRRWKPSEMKLDPFQEVVLESSSVDELREK DTYLDIPLVIRPYGSSQAFASVEEALHAFIQPEILDGPNQYFCERCKKKCDARKGLRFLHFPYLLTLQLKRFDFDYTTMH EAVEMAYKMMDLEEVIPLDCCRLVKYDEFHDYLERSYEGEEDTPMGLLLGGVKSTYMFDLLLETRKPDQVFQSYKPG FVESSETLDYQMAFADSHLWKLLDRHANTIRLFVLLPEQSPVSYSKRTAYQKAGGDSGNVDDDCERVKGPVGSLKSV EAILEESTEKLKSLSLQQQQDGDNGDSSKSTETSDFENIESPLNERDSSASVDNRELEQHIQTSDPENFQSEERSDSD YMLIYRLKDPARNAKFLEVGEYPEHIKNLVQKERELEEQEKROREIERNTCKIKLFCLHPTKQVMMENKLEVHKDKTLK LSEISGIPLDDIEFAKGRGTFPCDISVLDIHQDLDWNPKVSTLNVWPLYICDDGAVIFYRDKTEELMELTDEQRNELMKK EVMVKVHVVDLKAESVAAPITVRAYLNQTVTEFKQLISKAIHLPAETMRIVLERCYNDLRLLSVSSKTLKAEGFFRSNKV SSFSDDNKITIRLGRALKKGEYRVKVYQLLVNEQEPCKFLLDAVFAKGMTVRQSKEELIPQLREQCGLELSIDRFRLRK RIKLNDRMTFPEELDMSTFIDVEDEKSPQTESCTDSGAENEGSCHSDQMSNDFSNDDGVDEGICLETNSGTEKISKS RGEMQYMYFKAEPYAADEGSGEGHKWLMVHVDKRITLAAFKQHLEPFVGVLSSHFKVFRVYASNQEFESVRLNETL **ESSRLQKTGHRVTYSPRKEKALKIYLDGAPNKDLTQD** 

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>SGPr480 SEQID 73

**NLIVGLVLLTRGKDEEKAKYIFSLFSSESGNYVIREEMERMLHVVDGKVPDTLRKCFSEGEKVNYEKFRNWLFLNKDAF** PSPNAPLKRVLAYTGCFSRMQTIKEIHEYLSQRLRIKEEDMRLWLYNSENYLTLLDDEDHKLEYLKIQDEQHLVIEVRNK KRTPQLIHGRDYEMVPEPVWRALYHWYGANLALPRPVIKNSKTDIPELELFPRYLLFLRQQPATRTQQSNIWVNMGNV DMSWPEEMSFIANSSKIDRHKVPTEKGATGLSNLGNTCFMNSSIQCVSNTQPLTQYFISGRHLYELNRTNPIGMKGHM MGAKESRIGFLSYEEALRRVTDVELKRLKDAFKRTCGLSYYMGQHCFIREVLGDGVPPKVAEVIYCSFGGTSKGLHFN SHNWFIISMQWWQQWKEYVKYDANPVVIEPSSVLNGGKYSFGTAAHPMEQVEDRIGSSLSYVNTTEEKFSDNISTAS EASETAGSGFLYSATPGADVCFARQHNTSDNNNQCLLGANGNILLHLNPQKPGAIDNQPLVTQEPVKATSLTLEGGRI DWEVAAEAWDNHLRRNRSIVVDLFHGQLRSQVKCKTCGHISVRFDPFNFLSLPLPMDSYMHLEITVIKLDGTTPVRYG SPSTNEMFTLTTNGDLPRPIFIPNGMPNTVVPCGTEKNFTNGMVNGHMPSLPDSPFTGYIIAVHRKMMRTELYFLSSQ KNRPSLFGMPLIVPCTVHTRKKDLYDAVWIQVSRLASPLPPQEASNHAQDCDDSMGYQYPFTLRVVQKDGNSCAWC LRLNMDEKYTGLKKQLSDLCGLNSEQILLAEVHGSNIKNFPQDNQKVRLSVSGFLCAFEIPVPVSPISASSPTQTDFSS PWYRFCRGCKIDCGEDRAFIGNAYIAVDWDPTALHLRYQTSQERVVDEHESVEQSRRAQAEPINLDSCLRAFTSEEEI AKCYGDLVQELWSGTQKNVAPLKLRWTIAKYAPRFNGFQQQDSQELLAFLLDGLHEDLNRVHEKPYVELKDSDGRP FFSRWLLSGGVYVTLTDDSDTPTFYQTLAGVTHLEESDIIDLEKRYWLLKAQSRTGRFDLETFGPLVSPPIRPSLSEGL FNAFDENRDNHIDFKEISCGLSACCRGPLAERQKFCFKVFDVDRDGVLSRVELRDMVVALLEVWKDNRTDDIPELHM DLSDIVEGILNAHDTTKMGHLTLEDYQIWSVKNVLANEFLNLLFQVCHIVLGLRPATPEEEGQIIRGWLERESRYGLQA

KGRLRLPQIGSKNKLSSSKENLDASKENGAGQICELADALSRGHVLGGSQPELVTPQDHEVALANGFLYEHEACGNG YSNGQLGNHSEEDSTDDQREDTRIKPIYNLYAISCHSGILGGGHYVTYAKNPNCKWYCYNDSSCKELHPDEIDTDSAY QGDELSEPRILAREVKKVDAQSSAGEEDVLLSKSPSSLSANIISSPKGSPSSSRKSGTSCPSSKNSSPNSSPRTLGRS GENEMYYCSKCKTHCLATKKLDLWRLPPILIIHLKRFQFVNGRWIKSQKIVKFPRESFDPSAFLVPRDPALCQHKPLTP LFYEQQGIDYAQFLPKTDGKKMADTSSMDEDFESDYKKYCVLQ

>SGPr431 SEQID 74

SETSLOEVASKAAVLTETPRTSDGEKTLIEKMFGGKLRTHIRCLNCRSTSQKVEAFTDLSLAFCPSSSLENMSVQDPAS MDKILEGLVSSSHPLPLKRVIVRKVVESAEHWLDEAQCEAMFDLTTRLILEGQDPFQRQVGHQVLEAYARYHRPEFES PKGKLSITFCQQLVRTIGHFQCVSTQERELREYVSQVTKVSNLLQNIWKAEPATLLPSLQEVFASISSTDASFEPSVALA NGCNSLMKKLQHLFAFLAHTQREAYAPRIFFEASRPPWFTPRSQQDCSEYLRFLLDRLHEEEKILKVQASHKPSEILEC TPSVTDLLNYFLAPEILTGDNQYYCENCASLQNAEKTMQITEEPEYLILTLLRFSYDQKYHVRRKILDNVSLPLVLELPVK YHQSEALALASSQSHLLGRDSPSAVFEQDLENKEMSKEWFLFNDSRVTFTSFQSVQKITSRFPKDTAYVLLYKKQHST FFNKTFVLGLLHQGYHSLDRKDVAILDYIHNGLKLIMSCPSVLDLFSLLQVEVLRMVCERPEPQLCARLSDLLTDFVQCI RITSFSSLSESWSVDVDFTDLSENLAKKLKPSGTDEASCTKLVPYLLSSVVVHSGISSESGHYYSYARNITSTDSSYQM NGLSGNNPTSGLWINGDPPLQKELMDAITKDNKLYLQEQELNARARALQAASASCSFRPNGFDDNDPPGSCGPTGG RLWFPLVRPGALAVLSHMLLSFQHSPEAFHLIVPHVVNLVHSFKNDGLPSSTAFLVQLTELIHCMMYHYSGFPDLYEPI EAIKDFPKPSEEKIKLILNQSAWTSQSNSLASCLSRLSGKSETGKTGLINLGNTCYMNSVIQALFMATDFRRQVLSLNI SLVQHIPLQMITVLIRSLTTDPNVKDASMTQALCRMIDWLSWPLAQHVDTWVIALLKGLAAVQKFTILIDVTLLKIELVFN SPSIQDGGLMQASVPGPSEEPVVYNPTTAAFICDSLVNEKTIGSPPNEFYCSENTSVPNESNKILVNKDVPQKPGGET GGGGFNTVGRLVF

FIG. 2G

>SGPr429 SEQID 75

PSKSQTRKPKCGKGTHCSRNAYMLVYRLQTQEKPNTTVQVPAFLQELVDRDNSKFEEWCIEMAEMRKQSVDKGKAK HEEVKELYQRLPAGAEPYEFVSLEWLQKWLDESTPTKPIDNHACLCSHDKLHPDKISIMKRISEYAADIFYSRYGGGPR SKFYELELNIQGHKQLTDCISEFLKEEKLEGDNRYFCENCOSKQNATRKIRLLSLPCTLNLQLMRFVFDRQTGHKKKLN YIGFSEILDMEPYVEHKGGSYVYELSAVLIHRGVSAYSGHYIAHVKDPQSGEWYKFNDEDIEKMEGKKLQLGIEEDLE MAPRLQLEKAAWRWAETVRPEEVSQEHIETAYRIWLEPCIRGVCRRNCKGNPNCLVGIGEHIWLGEIDENSFHNIDDP NCERRKKNSFVGLTNLGATCYVNTFLQVWFLNLELRQALYLCPSTCSDYMLGDGIQEEKDYEPQTICEHLQYLFALLQ .TVKALCKECVVERCRILRLKNQLNEDYKTVNNLLKAAVKGDGFWVGKSSLRSWRQLALEQLDEQDGDAEQSNGKM NSNRRYIDPSGFVKALGLDTGQQQDAQEFSKLFMSLLEDTLSKQKNPDVRNIVQQQFCGEYAYVTVCNQCGRESKLL **VGSTLNKDESKEERKEEEELNFNEDILCPHGELCISENERRLVSKEAWSKLQQYFPKAPEFPSYKECCSQCKILEREG** 

GLMFTFASMTKEDSKLIALIWPSEWQMIQKLFVVDHVIKITRIEVGDVNPSETQYISEPKLCPECREGLLCQQQRDLREY RHRKVRGEKALLVSANQTLKELKIQIMHAFSVAPFDQNLSIDGKILSDDCATLGTLGVIPESVILLKADEPIADYAAMDDV EENEALHKMIANEQKTSLPNLFQDKNRPCLSNWPEDTDVLYIVSQFFVEEWRKFVRKPTRCSPVSSVGNSALLCPHG TOATIYVHKVVDNKKVMKDSAPELNVSSSETEEDKEEAKPDGEKDPDFNQSNGGTKRQKISHQNYIAYQKQVIRRSM MOVCMPEEGFKGTGLLGH

OSSSELS OSESOL

>SGPr503 SEQID 76

MNSLLQGLSACPAFIRWLEEFTSQYSRDQKEPPSHQYLSLTLLHLLKALSCQEVTDDEVLDASCLLDVLRMYRWQISS PVLNQPGAPKTQIFMNGACSPSLLPTLSAPMPFPLPVVPDYSSSTYLFRLMAVVVHHGDMHSGHFVTYRRSPPSARN CKHCEHQSPVRFDTFDSLSLSIPAATWGHPLTLDHCLHHFISSESVRDVVCDNCTKIEAKGTLNGEKVEHQRTTFVKQ MLSSRAEAAMTAADRAIQRFLRTGAAVRYKVMKNWGVIGGIAAALAAGIYVIWGPITERKKRRKGLVPGLVNLGNTCF -KLGKLPQCLCIHLQRLSWSSHGTPLKRHEHVQFNEFLMMDIYKYHLLGHKPSQHNPKLNKNPGPTLELQDGPGAPT FEEQDAHELFHVITSSLEDERDRQPRVTHLFDVHSLEQQSEITPKQITCRTRGSPHPTSNHWKSQHPFHGRLTSNMV PLSTSNQWLWVSDDTVRKASLQEVLSSSAYLLFYERVLSRMQHQSQECKSEE

>SGPr427 SEQID 77

-REDEVNTRGAYILFYQKRNSIPPWSASSSMRGSTSSSLSDHWLLRLGSHAGSTRGSLLSWSSAPCPSLPQVPDSPIF SQSIVSLLTGTAGEDEKSASPRSNVALPANSEDGGRAIERGPAGVPCPSAQPNHCLAPGNSDGPNTARKLKENAGQD RALWTREYTPQLSAEFKNAVSKYGSQFQGNSQHDALEFLLWLLDRVHEDLEGSSRGPVSEKLPPEATKTSENCLSPS PHVAQRSTSPEAGLGPWPSWKQPDCLPTSYPLDFLYDLYAVCNHHGNLQGGHYTAYCRNSLDGQWYSYDDSTVEP MDLGPGDAAGGGPLAPRPRRRRSLRRLFSRFLLALGSRSRPGDSPPRPQPGHCDGDGEGGFACAPGPVPAAPGSP SCTLDECFQFYTKEEQLAQDDAWKCPHCQVLQQGMVKLSLWTLPDILIIHLKRFCQVGERRNKLSTLVKFPLSGLNMA GEERPPGPQPQLQLPAGDGARPPGAQGLKNHGNTCFMNAVVQCLSNTDLLAEFLALGRYRAAPGRAEVTEQLAALV alrkmvaeeggvpadevilvelypsgfqrsffdeedlntiaegdnvyafqvppspsqgtlsahplglsasprlaareg QRFSLSLHSESKVLILFCNLVGSGQQASRFGPPFLIREDRAVSWAQLQQSILSKVRHLMKSEAPVQNLGSLFSIRVVGL **AQLPLGQSFVQSHFQAQYRSSLTCPHCLKQSNTFDPFLCVSLPIPLRQTRFLSVTLVFPSKSQRFLRVGLAVPILSTVA** SVACSYLSPKDSRPLCHWAVDRVLHLRRPGGPPHVKLAVEWDSSVKERLFGSLQEERAQDADSVWQQQQAHQQH KLPRKFDLPLTVMPSVEHEKPARPEGQKAMNWKESFQMGSKSSPPSPYMGFSGNSKDSRRGTSELDRPLQGTLTL RSVFRKKENRRNERAEVSPQVPPVSLVSGGLSPAMDGQAPGSPPALRIPEGLARGLGSRLERDVWSAPSSLRLPR INSLCNQEKGGLEPRRLVRGVKGRSISMKAPTTSRAKQGPFKTMPLRWSFGSKEKPPGASVELVEYLESRRRPRST *<b>KASRAPRGSALGMSQRTVPGEQASYGTFQRVKYHTLSLGRKKTLPESSI* 

>SGPr092\_SEQID 78

WAHEDGWPFDGPGGILGHAFLPNSGNPGVVHFDKNEHWSASDTGYNLFLVATHEIGHSLGLQHSGNQSSIMYPTYW HQPHCGVPDGSDTSISPGRCKWNKHTLTYRIINYPHDMKPSAVKDSIYNAVSIWSNVTPLIFQQVQNGDADIKVSFWQ MQLVILRVTIFLPWCFAVPVPPAADHKGWDFVEGYFHQFFLTKKESPLLTQETQTQLLQQFHRNGTDLLDMQMHALI YHDPRTFQLSADDIQRIQHLYGEKCSSDIP

SGPr359 SEQID 79

RHVQQIDAAVYLREPQKTLFFVGDEYYSYDERKRKMEKDYPKNTEEEFSGVNGQIDAAVELNGYIYFFSGPKTYKYDT FGLQVTGKLDQTTMNVIKKPRCGVPDVANYRLFPGEPKWKKNTLTYRISKYTPSMSSVEVDKAVEMALQAWSSAVPL GLAHSTDPSALMYPTYKYKNPYGFHLPKDDVKGIQALYGPRKVFLGKPTLPHAPHHKPSIPDLCDSSSSFDAVTMLGK SFVRINSGEADIMISFENGDHGDSYPFDGPRGTLAHAFAPGEGLGGDTHFDNPEKWTMGTNGFNLFTVAAHEFGHAL ELLLFKDRIFWRRQVHLRTGIRPSTITSSFPQLMSNVDAAYEVAERGTAYFFKGPHYWITRGFQMQGPPRTIYDFGFP MKVLPASĞLAVFLIMALKFSTAAPSLVAASPRTWRNNYRLAQAYLDKYYTNKEGHQIGEMVARGSNSMIRKIKELQAF **EKEDVVSVVKSSSWIGC** 

>SGPr104 1 SEQID 80

MNVALQELGAGSNMVEYKRATLRDEDAPETPVEGGASPDAMEVGFQKGTRQLLGSRTQLELVLAGASLLLAALLLGC -KHLLENTTFNSSSEAEQKTQRFYLSCLQVERIEELGAQPLRDLIEKIGGWNITGPWDQDNFMEVLKAVAGTYRATPFF WQNESLAAFRNHTACMEEQYNQYQVNGERLNGRQTLGENIADNGGLKAAYNAYKAWLRKHGEEQQLPAVGLTNHQ EEALGQLVWMDEKTRQAAKEKADAIYDMIGFPDFILEPKELDDVYDGYEISEDSFFQNMLNLYNFSAKVMADQLRKPP LVALGVQYHRDPSHSTCLTEACIRVAGKILESLDRGVSPCEDFYQFSCGGWIRRNPLPDGRSRWNTFNSLWDQNQAI TVYISADSKSSNSNVIQVDQSGLFLPSRDYYLNRTANEKVLTAYLDYMEELGMLLGGRPTSTREQMQQVLELEIQLANI NLVQKTTSSLDRRFESAQEKLLETLYGTKKSCVPRWQTCISNTDDALGFALGSLFVKATFDRQSKEIAEGMISEIRTAF TVPQDQRRDEEKIYHKMSISELQALAPSMDWLEFLSFLLSPLELSDSEPVVVYGMDYLQQVSELINRTEPSILNNYLIW SRDQWSMTPQTVNAYYLPTKNEIVFPAGILQAPFYARNHPKALNFGGIGVVMGHELTHAFDDQGREYDKEGNLRPW LFFVGFAQVWCSVRTPESSHEGLVTDPHSPARFRVLGTLSNSRDFLRHFGCPVGSPMNPGQLCEVW

>SGPr303\_SEQID\_81

EDEKVTLSFPSTLQTGTGTLKIDFVGELNDKMKGFYRSKYTTPSGEVRYAAVTQFEATDARRAFPCWDERAIKATFDIS MPEKRPFERLPADVSPINCSLCLKPDLLDFTFEGKLEAAAQVRQATNQIVMNCADIDIITASYAPEGDEEIHATGFNYQN ALEVAAKTLPFYNDYFNVPYPLPKIDLIAIADFAAGAMENWDLVTYRETALLIDPKNSCSSSRQWVALVVGHELAHQWF LVVPKDRVALSNMNVIDRKPYPDDENLVEVKFARTPVTSTYLVAFVVGEYDFVETRSKDGVCVCVYTPVGKAEQGKF

GNLVTMEWWTHLWLNEGFASWIEYLCVDHCFPEYDIWTQFVSADYTRAQELDALDNSHPIEVSVGHPSEVDEIFDAIS YSKGASVIRMLHDYIGDKVKKKTLSI

DOMODOLO LOGICA

>SGPr402 1 SEQID 82

/VQQSLTPHWGHHLHLKKNPKVQWFQQQTLQRRVKRSVVVPTDPWFSKQWYMNSEAQPDLSILQAWSQGLSGQGI VVSVLDDGIEKDHPDLWANYDPLASYDFNDYDPDPQPRYTPSKENRHGTRCAGEVAAMANNGFCGVGVAFNARIGG LGQLCLAYCPPRFFNHTRLVTAGPGHTAAPALRVCSSCHASCYTCRGGSPRDCTSCPPSSTLDQQQGSCMGPTTPD WRPAPIALWLRLVLALALVRPRAVGWAPVRAPIYVSSWAVQVSQGNREVERLARKFGFVNLGPIFPDGQYFHLRHRG VRMLDGTITDVIEAQSLSLQPQHIHIYSASWGPEDDGRTVDGPGILTREAFRRGVTKGRGGLGTLFIWASGNGGLHYD **ALALEANPFLTWRDMQHLVVRASKPAHLQAEDWRTNGVGRQVSHHYGYGLLDAGLLVDTARTWLPTQPQRKCAVR** MSTHFWDENPQGVWTLGLENKGYYFNTGTLYRYTLLLYGTAEDMTARPTGPQVTSSACVQRDTEGLCQACDGPAYI VQSRPTPILPLIYIRENVSACAGLHNSIRSLEHVQAQLTLSYSRRGDLEISLTSPMGTRSTLVAIRPLDVSTEGYNNWVF NCNCDGYTNSIHTLSVGSTTQQGRVPWYSEACASTLTTTYSSGVATDPQIVTTDLHHGCTDQHTGTSASAPLAAGMI SRPRLRAAACPHHRCPASAMVLSLLAVTLGGPVLCGMSMDLPLYAWLSRARATPTKPQVWLPAGT

>SGPr434 SEQID 83

CATNYGENLCYGDSGGPLACEVEGRWILAGVLSWEKACVKAQNPGVYTRITKYTKWIKKQMSNGAFSGPCASACLLF GPGRQGGCAGRRSTALPLRAPLRARRPGPRSERMGAATCRGSRIPSGPPVQGERSAPRFGVTSLSLWPADFKDNW RIAGSRQEVALAGEPADQQQTHLRRLPYRQTLGYKEDTTNPVCGEPWWSEDLEMTRHWPWEVSLRMENEHVCGG QPICLPEPNFNLKVGTQCWVTGWSQVKQRFSGSTANSMLTPELQEAEVFIMDNKRCDRHYKKSFFPPVVPLVLGDMI alidpswyytaahcsqgtkeysyylgtsklqpmnfsralwypyrdiimhpkywgrafimgdvalyhlqtpytfseyy

>SGPr446 1 SEQID 84

VVSVQRAFVHPKFSTVTTIRNDLALLQLQHPVNFTSNIQPICIPQENFQVEGRTRCWVTGWGKTPERGEKLASEILQDV I\_TPVCGRTPLRIVGGVDAEEGRWPWQVSVRTKGRHICGGTLVTATWVLTAGHCISSRFHYSVKMGDRSVYNENTSV DQYIMCYEECNKIIQKALSSTKDVIIKGMVCGYKEQGKDSCQGDSGGRLACEYNDTWVQVGIVSWGIGCGR

>SGPr447 SEQID 85

QKHYYPSEWTVQLGELTSRPTPWNLRAYSSRYKVQDIIVNPDALGVLRNDIALLRLASSVTYNAYIQPICIESSTFNFVH MGARGAL\_LALLLARAGLGKPEACGHREIHALVAGGVESARGRWPWQASLRLRRRHRCGGSLLSRRWVLSAAHCF

## CORDERIN CONTOL

RPDCWVTGWGLISPSGTPLPPPYNLREAQVTILNNTRCNYLFEQPSSRSMIWDSMFCAGAEDGSVDTCKGDSGGPI VCDKDGLWYQVGIVSWGMDCGQPNRPGVYTNISVYFHWIRRVMSHSTPRPNPSQLLLLLALLWAP

-PGMVCTSAVGELPSCEGLSGAPLVHEVRGTWFLAGLHSFGDACQGPARPAVFTALPAYEDWVSSLDWQVYFAEEP YDMALLLLAQPVTLGASLRPLCLPYPDHHLPDGERGWVLGRARPGAGISSLQTVPVTLLGPRACSRLHAAPGGDGSPI GASCWATGWDQDTSDAPGTLRNLRLISRPTCNCIYNQLHQRHLSNPARPGMLCGGPQPGVQGPCQGDSGGPVL NQSNSIFGCIFYTLQLLLGLQAAQRACGQRGPGPPKPQEGNTVPGEWPWQASVRRQGAHICSGSLVADTWVLTAAH CFEKAAATELNSWSVVLGSLQREGLSPGAEEVGVAALQLPRAYNHYSQGSDLALLQLAHPTTHTPLCLPQPAHRFPF CLEPDGHWVQAGIISFASSCAQEDAPVLLTNTAAHSSWLQARVQGAAFLAQSPETPEMSDEDSCVACGSLRTAGPQ AGAPSPWPWEARLMHQGQLACGGALVSEEAVLTAAHCFIGRQAPEEWSVGLGTRPEEWGLKQLILHGAYTHPEGG >SGPr432\_1\_SEQID\_86 MGSTWGSPGWVRLALCLTGLVLSLYALHVKAARARDRDYRALCDVGTAISCSRVFSSRWGRGFGLVEHVLGQDSII EPEAEPGSCLANISQPTSC

>SGPr529\_SEQID\_87

DQSWVLTAAHCGNKPLWARVGDDHLLLLQGEQLRRTTRSVVHPKYHQGSGPILPRRTDEHDLMLLKLARPVVPGPR WRAPHLHLSAASGARALAKLLPLLMAQLWAAEAALLPQNDTRLDPEAYGAPCARGSQPWQVSLFNGLSFHCAGVLV VRALQLPYRCAQPGDQCQVAGWGTTAARRVKYNKGLTCSSITILSPKECEVFYPGVVTNNMICAGLDRGQDPCQSD SGGPLVCDETLQGILSWGVYPCGSAQHPAVYTQICKYMSWINKVIRSN

QYDDLNMHLEKLRVVQISRKECAKRVNQLSRNMICAWNEPGTNGQGPGEVGGPLVCQKKNKSTWYQLGIISWGVGC NVQCGHRPAFPNSSWLPFHERLQVQNGECPWQVSIQMSRKHLCGGSILHWWWVLTAAHCFRRTLLDMAVVNVTVV MGTRTFSNIHSERKQVQKVIIHKYYKPPQLDSDLSLLLLATPVQFSNFKMPVCLQEEERTWDWCWMAQWVTTNGYD SQKNMPGVYTELSNYLLWIERKTVLAGKPYKYEPDSVYALLLSPWAILLLYFVMLLLS

SGPr425 SEQID 89

EDYLSYETVFENGTRTLTRVKVQDLVLEPTQNITTKGVSVRRKRQVYGTDSRFSILDKRFLTNFPFSTAVKLSTGCSGIL MENMLLWLIFFTPGWTLIDGSEMEWDFMWHLRKVPRIVSERTFHLTSPAFEADAKMMVNTVCGIECQKELPTPSLSEL SPOHVLTAAHCVHDGKDYVKGSKKLRVGLLKMRNKSGGKKRRGSKRSRREASGGDQREGTREHLPERAKGGRRR KKSGRGQRIAEGRPSFQWTRVKNTHIPKGWARGGMGDATLDYDYALLELKRAHKKKYMELGISPTIKKMPGGMIHFS

## OSESELE . OEESIL

GFDNDRADQLVYRFCSVSDESNDLLYQYCDAESGSTGSGVYLRLKDPDKKNWKRKIIAVYSGHQWVDVHGVQKDYN VAVRITPLKYAQICLWIHGNDANCAYG

>SGPr548 SEQID 90

TTSRVIPHPRYEARSHRNDIMLLRLVQPARLNPQVRPAVLPTRCPHPGEACVVSGWGLVSHNEPGTAGSPRSQVSLP DTLHCANISIISDTSCDKSYPGRLTNTMVCAGAEGRGAESCEGDSGGPLVCGGILQGIVSWGDVPCDNTTKPGVYTKV MGDPEGSAEWGWGKGIPVVRRNLLTVDGISLCLEGSWWRQKGPASPGFSHSLPRLQPNPGPSSTMWLLLTLSFLLA STAAQDGDKLLEGDECAPHSQPWQVALYERGRFNCGASLISPHWVLSAAHCQSRFMRVRLGEHNLRKRDGPEQLR CHYLEWIRETMKRN

>SGPr396 SEQID 91

MGPAGCAFTLLLLLGISVCGQPVYSSRVVGGQDAAAGRWPWQVSLHFDHNFICGGSLVSERLILTAAHCIQPTWTTFS **VSWGLECGKSLPGVYTNVIYYQKWINATISRANNLDFSDFLFPIVLLSLALLRPSCAFGPNTIHRVGTVAEAVACIQGWE** YTVWLGSITVGDSRKRVKYYVSKIVIHPKYQDTTADVALLKLSSQVTFTSAILPICLPSVTKQLAIPPFCWVTGWGKVKE SSDRDYHSALQEAEVPIIDRQACEQLYNPIGIFLPALEPVIKEDKICAGDTQNMKDSCKGDSGGPLSCHIDGVWIQTGV ENAWRFSPRGR

>SGPr426 SEQID 92

KPSFRLTRCGIRMTSSNMPLPASSSTQRIVQGRETAMEGEWPWQASLQLIGSGHQCGASLISNTWLLTAAHCFWKNK vddgpiqntlrqarvetistdvcnrkdvydglitpgmlcagfmegkidackgdsggplvydnhdiwyivgivswgqsc refiershqiermmsrifrhssvggrfikshviklspdeqgvdilivlifrypstdsaeqikkkiekalyqslktkqlsltin MMYAPVEFSEAEFSRAEYQRKQQFWDSVRLALFTLAIVAIIGIAIGIVTHFVVEDDKSFYYLASFKVTNIKYKENYGIRSS **DPTQWIATFGATITPPAVKRNVRKIILHENYHRETNENDIALVQLSTGVEFSNIVQRVCLPDSSIKLPPKTSVFVTGFGSI** ALPKKPGVYTRVTKYRDWIASKTGM

>SGPr552 SEQID 93

RIAEGLDAEEGEWPWQASLPQNNVYRRGATWLSNSWLITAAHCFIRVHDPKEWNVILSNPQTQSNIKNVIIQENYHYP AHDNDIAVVHLSSPVLYTSNIQKACLPDVNYIFLYNSEAVVTAWGSFKPLRTTSNVLHKGLVKIIDNRTCNNGEADGRVI TSGMLCAGFLEPRVDACQGDSGGPLVGTDSKGILAKGSLLVLKAGVNERALPNKPSVYTQVTYY

>SGPr405 SEQID 94

MVSKGGVAAEPEPHYCEDSERGPNTLTGPGSLPRGGGIEVGMEFPGCSGEGCVKPHEEAAREGAGRGKRAVPGPK **WPLSLWEATVKVRSNVLCNQTCRRRFPSNHTERFERLIKDDMLCAGDGNHGSWPGDNGGPLLCRRNCTWVQVEVV** 3KMAEFNWSMAFKGPAAGHEERLNSVSSRAKKGIGWDVAAASLRGVDHFSDLPPPLQVREELEACAFRVQVGQLRL <u> MRTRACERMYHKGPTAHGQVTIIKAAMPCAGRKGQGSCQAALRTEDL†PTTPNTEVSPRADPRLSQPEDIWPEWAW</u> RRQQGSAEGPAAGWTLEQETRGDVLEDKNERADEEILRLAPGKGRLPIDSKHLKPVISSFPVRSQELGEGAGAGTLR PETSWQGANHADAQRPAGRVPTMQRPRDMGQGQEWVCRPFTHVTCYPTAIPRPFTHVTCYLMAVPSTLTHVTCYP PVVVGTTMLLLLLFLAVSSLGSCSTGSPAPVPENDLVGIVGGHNTPGEVVVAVGADRRSLHFPEGHRPVHLPDSHQG SNADCERQTYDAFPGAGDRKFIQDDMICAGRTGRRTWKGDSGGPLVCKKKGTWLQAGVVSWGFYSDRPSIGVYTR YEDDQRTKVVEIVRHPQYNESLSAQGGADIALLKLEAPVPLSELIHPVSLPSASLDVPSGKTCWVTGWGVIGRGELLP **FAVPRPFTHVTCYLMAVPSTLTHITCYMMAVPRPFTHITCYPMAVPSTLTHVTCHPTAIPRPFTHITCYTMAIPRPSTTP** SVSVRGPGAAECQPDRRPPNYSVFFLGADIALLKLATSSLEFTDSDNCWNTGWGMVGLLDMLPPPYRPQQVKVLTL SWGKLCGLRGYPGMYTRVTSYVSWIRQPCPSAQTPAVVRRFVLPPNPDVEALTPSVMGSGAPLPPAPDLQEAEVPI PATRRPSPAPSPTSPATRWPSPGPSPMSPATR

>SGPr485\_1\_SEQID\_95

.LLLASPIKLDDLKVPICLPTQPGPATWRECWVAGWGQTNAADKNSVKTDLMKAPMVIMDWEECSKMFPKLTKNMLC AGYKNESYDACKGDSGGPLVCTPEPGEKWYQVGIISWGKSCGEKNTPGIYTSLVNYNLWIEKVTQLEGRPFNAEKRR \*PWQVSIQVRSEPFCGGSILNKWWILTAAHCLYSEELFPEELSVVLGTNDLTSPSMEIKEVASIILHKDFKRANMDNDIA <u>MLLFSVLLLLSLVTRTQLGPRTPLPEAGVAILGRARGAHRPQPPHPPSPVSECGDRSIFEGRTRYSRITGGMEAEVGE</u> SVKQKPMGSPVSGVPEPGSPRSWLLLCPLSHVLFRAILY

>SGPr534\_SEQID\_96

3VRTSDVVVAGEFDQGSDEENIQVLKIAKVFKNPKFSILTVNNDITLLKLATPARFSQTVSAVCLPSADDDFPAGTLCAT 1GWGKTKYNANKTPDKLQQAALPLLSNAECKKSWGRRITDVMICAGASGVSSCMGDSGGPLVCQKDGAWTLVGIVS MASLWLLSCFSLVGAAFGCGVPAIHPVLSGLSRIVNGEDAVPGSWPWQVSLQDKTGFHFCGGSLISEDWVVTAAHC *WGSRTCSTTTPAVYARVTKLIPWVQKILAAN* 

>SGPr390 SEQID 97

NPPEGRNSEHIRTSARTNSGHTIFKKCNTQPFLSTQGFHVDHTAELRGIRWTSSLRRETSDYHRTLTPTLEALLHFLLR MEPTVADVHLVPRTTKEVPALDAACCRAASIGVVATSLVVLTLGVLLGGMNNSRHAALRAATLPGKVYSVTPEASKT

**DPTKWVAYVGATYLSGSEASTVRAQVVQIVKHPLYNADTADFDVAVLELTSPLPFGRHIQPVCLPAATHIFPPSKKCLIS** GWGNTQEGNATKPELLQKASVGIIDQKTCSVLYNFSLTDRMICAGFLEGKVDSCQGDSGGPLACEEAPGVFYLAGIVS GWGYLKEDFRKHLPRPAMVKPEVLQKATVELLDQALCASLYGHSLTDRMVCAGYLDGKVDSCQGDSGGPLVCEEPS WGIGCAQVKKPGVYTRITRLKGWILEIMSSQPLPMSPPSTTRMLATTSPRTTAGLTVPGATPSRPTPGAASRVTGQPA NSTLSAVSTTARGQTPFPDAPEATTHTQLPDCGLAPAALTRIVGGSAAGRGEWPWQVSLWLRRREHRCGAVLVAER DDQEDCSDGSDEAHCECGLQPAWRMAGRIVGGMEASPGEFPWQASLRENKEHFCGAAIINARWLVSAAHCFNEFQ WLLSAAHCFDVYGDPKQWAAFLGTPFLSGAEGQLERVARIYKHPFYNLYTLDYDVALLELAGPVRRSRLVRPICLPEP PLQTLSLGLEEELLQRGIRARLREHGISLAAYGTIVSAELTGRHKGPLAERDFKSGRCPGNSFSCGNSQCVTKVNPEC GRFFLAGIVSWGIGCAEARRPGVYARVTRLRDWILEATTKASMPLAPTMAPAPAAPSTAWPTSPESPVVSTPTKSMQ ALSTVPLDWVTVPKLQECGARPAMEKPTRVVGGFGAASGEVPWQVSLKEGSRHFCGATVVGDRWLLSAAHCFNHT KVEQVRAHLGTASLLGLGGSPVKIGLRRVVLHPLYNPGILDFDLAVLELASPLAFNKYIQPVCLPLAIQKFPVGRKCMIS APRPPDGTRCVITGWGSVREGGSMARQLQKAAVRLLSEQTCRRFYPVQISSRMLCAGFPQGGVDSCSGDAGGPLA CREPSGRWVLTGVTSWGYGCGRPHFPGVYTRVAAVRGWIGQHIQE

COSSOCIE CENSOL

>SGPr521 SEQID 98

MARSLLLPLQILLLSLALETAGEEAQGDKIIDGAPCARGSHPWQVALLSGNQLHCGGVLVNERWVLTAAHCKMNEYTV HLGSDTLGDRRAQRIKASKSFRHPGYSTQTHVNDLMLVKLNSQARLSSMVKKVRLPSRCEPPGTTCTVSGWGTTTS PDVTFPSDLMCVDVKLISPQDCTKVYKDLLENSMLCAGIPDSKKNACNGDSGGPLVCRGTLQGLVSWGTFPCGQPN **DPGVYTQVCKFTKWINDTMKKHR** 

SGPr530\_1\_SEQID\_99

HQTQHTQKMSVHRIITHPDFEKLHPFGSDIAMLQLHLPMNFTSYIVPVCLPSRDMQLPSNVSCWITGWGMLTEDHKRV QLSPPFYLQEGKVGLIENTLCNTLYGQRTAKARPKLCTRRCCVGGYFSTGKSICKGDSGGPLVCYLPSAWVLVGLAS VSTVCGKPKVVGKIYGGRDAAAGQWPWQASLLYWGSHLCGAVLIDSCWLVSTTHCFLNKSQAPKNYQVLLGNIQLY **WGLDCRHPAYPSIFTRVTYFINWIDEIMRLTPLSDPALAPH** 

>SGPr520\_SEQID\_100

PKFDPRTFHNDLALVQLWTPVSPGGSARPVCLPQEPQEPPAGTACAIAGWGALFEDGPEAEAVREARVPLLSTDTCR RALGPGLRPSTMLCAGYLAGGVDSCQGDSGGPLTCSEPGPRPREVLFGVTSWGDGCGEPGKPGVYTRVAVFKDWL QEQMSAASSSREPSCRELLAWDPPQELQADAARLCAFYARLCPGSQGACARLAHQQCLQRRRRCELRSLAHTLLGL MLLAVILLI\_PLPSSWFAHGHPLYTRLPPSALQVFTLLLGAETVLGRNLDYVCEGPCGERRPSTANVTRAHGRIVGGSA APPGAWPWLVRLQLGGQPLCGGVLVAASWVLTAAHCFVGCRSTRSAPNELLWTVTLAEGSRGEQAEEVPVNRILPH

## 

LRNAQELLGPRPGLRRLAPALALPAPALRESPLHPARELRLHSGSRAAGTRFPKRRPEPRGEANGCPGLEPLRQKLA ALQGAHAWILQVPSEHLAMNFHEVLADLGSKTLTGLFRAWVRAGLGGRHVAFSGLVGLEPATLARSLPRLLVQALQA FRVAALAEGEPEGPWMDVGQGPGLERKGHHPLNPQVPPARQP

MSPDIALLYLKHKVKFGNAVQPICLPDSDDKVEPGILCLSSGWGKISKTSEYSNVLQEMELPIMDDRACNTVLKSMNLP ?LGRTMLCAGFPDWGMDACQGDSGGPLVCRRGGGIWILAGITSWVAGCAGGSVPVRNNHVKASLGIFSKVSELMDF EICAVTGWGSISAELSLNVSSLDGGLASRLQQIQVHVLEREVCEHTYYSAHPGGITEKMICAGFAASGEKDFCQGDSG FMSPGPLVRVTFHALVRGAFGISYIVLKVLGPKDSKITRLSQSSNREHLVPCEDVLLTKPEGIMQIPRNSHRTTMGCQW TQNLFTGLDRGQPLSKVGSRYITKALSSVQEVNGSQRDKIILIKFTSLDMEKQVGCDHDYVSLRSSSGVLFSKVCGKIL \*SPLLAETSEAMVPFVSDTEDSGSGFELTVTAVQKSEAGSGCGSLAILVEEGTNHSAKYPDLYPSNTRCHWFICAPEK HIIKLTFEDFAVKFSPNCIYDAVVIYGDSEEKHKLAKLCGMLTITSIFSSSNMTVIYFKSDGKNRLQGFKARFTILPSESLN (FEPKLPPQNNPVSTVKAILHDVCGIPPFSPQWLSRRIAGGEEACPHCWPWQVGLRFLGDYQCGGAIINPVWILTAAH RLVAPLNHIIQLNIINFPMKPTTFVCHGHLRVYEGFGPGKKLIGRMLMSTELSWFLSQFSTKKTTASCGETAVSMKMMY SVQLKNNPLSWTIIAGDHDRNLKESTEQVRRAKHIIVHEDFNTLSYDSDIALIQLSSPLEYNSVVRPVCLPHSAEPLFSS <PRGFFPTPRYLLDYRGRLECSWVLRVSASSMAKFTIEYLSLLGSPVCQDSVLIIYEERHSKRKTAGGLHGRRLYSMT</pre> GPLVCRHENGPFVLYGIVSWGAGCVQPWKPGVFARVMIFLDWIQSKINGKLFSNVIKTITSFFRVGLGTVSCCSEAEL **TSIFLALQNTCYHALPHEVVLRIK** 

-HRGPAIPDWQRHNSHEQGRHPDLRQNLEAPVMSDRECQKTEQGKSHRNSLCVKFVKVFSRIFGEVAVATVICKDKL QTINPIQIVRYWNYSHSAPQDDLMLIKLAKPAMLNPKVQPLTLATTNVRPGTVCLLSGLDWSQENSGLWQLEPPGHLT >SGPr507\_2\_SEQID\_102 MKYVFYLGVLAGTFFFADSSVQKEDPAPYLVYLKSHFNPCVGVLIKPSWVLAPAHCYLPNLKVMLGNFKSRVRDGTE QGIEVGHFMGGDVGIYTNVYKYVSWIENTAKDK

>SGPr559 SEQID 103

KCIELIARCDGVSDCKDGEDEYRCVRVGGQNAVLQVFTAASWKTMCSDDWKGHYANVACAQLGFPSYVSSDNLRVS SLEGGFREEFVSIDHLLPDDKVTALHHSVYVREGCASGHVVTLQCTACGHRRGYSSRIVGGNIMSLLSQWPWQASLQ FQGYHLCGGSVITPLWIITAAHCVYDLYLPKSWTIQVGLVSLLDNPAPSHLVEKIVYHSKYKPKRLGNDIALMKLAGPLT FNEMIQPVCLPNSEENFPDGKVCWTSGWGATEDGAGDASPVLNHAAVPLISNKICNHRDVYGGIISPSMLCAGYLTG MGENDPPAVEAPFSFRSLFGLDDLKISPVAPDADAVAAQILSLLPLKFFPIIVIGIIALILALAIGLGIHFDCSGKYRCRSSF 3VDSCQGDSGGPLVCQERRLWKLVGATSFGIGCAEVNKPGVYTRVTSFLDWIHEQMERDLKT

>SGPr567 1 SEQID 104

PAQASPAQASPAQASPARASPALASLSRSSSGRSSSARSASVTTSPTRVYLVRATPVGAVPIRSSPARSAPATRATRE MERDSHGNASPARTPSAGASPAQASPAGTPPGRASPAQASPAQASPAGTPPGRASPAQASPAGAPAGTPPGRASPGRAS SPVQFWQGHTGIRYKEQRESCPKHAVRCDGVVDCKLKSDELGCVRFDWDKSLLKIYSGSSHQWLPICSSNWNDSYS EKTCQQLGFESAHRTTEVAHRDFANSFSILRYNSTIQESLHRSECPSQRYISLQCSHCGLRAMTGRIVGGALASDSKW >WQVSLHFGTTHICGGTLIDAQWVLTAAHCFFVTREKVLEGWKVYAGTSNLHQLPEAASIAEIIINSNYTDEEDDYDIAL <u> WRLSKPLTLSAHIHPACLPMHGQTFSLNETCWITGFGKTRETDDKTSPFLREVQVNLIDFKKCNDYLVYDSYLTPRMM</u> CAGDLRGGRDSCQGDSGGPLVCEQNNRWYLAGVTSWGTGCGQRNKPGVYTKVTEVLPWIYSKMESEVRFRKS

>SGPr479 1 SEQID 105

HYAGLHVCGGSILNEYWVLSAAHCFHRDKNIKIYDMYVGLVNLRVAGNHTQWYEVNRVILHPTYEMYHPIGGDVALVQ -KTRIVFSESVLPVCLATPEVNLTSANCWATGWGLVSKQGETSDELQEVQLPLILEPWCHLLYGHMSYIMPDMLCAGD <u>MAAPASVĪMGPLGPSĀLGLLLLLLVVAPPRVAALVHRQPENQGISLTGSVACGRPSMEGKILGGVPAPERKWPWQVSV</u> LNAKTVCEGDSGGPLVCEFNRSWLQIGIVSWGRGCSNPLYPGVYASVSYFSKWICDNIEITPTPAQPAPALSPALGPT .SVLMAMLAGWSVL

>SGPr489 1 SEQID 106

C G G D S G G S L M C R N K G A W D S G W S I W E A Q V G G S L E S R S S R P S L G N K V R L C L T N N F F K L A G C G T W C S E Q D V I V S G A EGKLHFPESLHLYYESKQRCVWTLLVPEEMHVLLSFSHLDVESCHHSYLSMYSLEDRPIGKFCGESLPSSILIGSNSLR 3DCTSDYVTVHSDVERKKEIARLCGYDVPTPVLSPSSIMLISFHSDENGTCRGFQAIVSFIPKAVYPDLNISISEDESMFL CGGSIVSPQWVITAAHCIANRNIVSTLNVTAGEYDLSQTDPGEQTLTIETVIIHPHFSTKKPMDYDIALLKMAGAFQFGH =VGPICLPELREQFEAGFICTTAGWGRLTEGGVLSQVLQEVNLPILTWEECVAALLTLKRPISGKTFLCTGFPDGGRDA MSLKMLISRNKLILLLGIVFFERGKSATLSLPKAPSCGQSLVKVQPWNYFNIFSRILGGSQVEKGSYPWQVSLKQRQKH KFVSDATDYAAGFNLTYKALKPNYIPGCSYLTVLFEEGLIQSLNYPENYSDKANCDWIFQASKHHLIKLSFQSLEIEES

>SGPr465\_1\_SEQID\_107

HADYNELHRMGSDITLLQLHHHVEFSSHILPACLPEPTTWLAPDSSCWISGWGMVTEDVFLPEPFQLQEAEVGVMDN IVCGSFFQPQYPGQPSSSDYTIHEDMLCAGDLITGKAICRVNSRGPLVCPLNGTWFLMGLSSWSLDCCSPVGPRVFT RWPWQASLLYLGGHICGAALIDSNWVASAAHCFQRCIFPPRAPLSTNPSDYRILLGYDQQSHPTEHSKQMTVNKIMV **RLPYFTNWISQKKRESTPPDPALAPPQETPPALDSMTSQGIVHKPGLCAALLAAHMFLLLLILLGSL** 

>SGPr524 1 SEQID 108

COMMONIT CONTOL

MDKENSDVSAAPADLKISNISVQVVSAQKKLPVRRPPLPGRRLPLPGRRPPQRPIGKAKPKKQSKKKVPFWNVQNKIIL <u>EYGSYNISOPCPVGSFRCSSGLCVPQAQRCDGVNDCFDESDELFCVSPQPACNTSSFRQHGPLICDGFRDCENGRD</u> VADVSSNNKGGLLVHFWIVFVMPRAKGHIFCEDCVAAILKDSIQTSIINRTSVGSLQGLAVDMDSVVLNGDCWSFLKK KRRENGAVSTDKGCSQYFYAEHLSLHYPLEISAASGRLMCHFKLVAIVGYLIRLSIKSIQIEADNCVTDSLTIYDSLLPIR /GSAYCGASVISREWLLSAAHCFHGNRLSDPTPWTAHLGMYVQGNAKFVSPVRRIVVHEYYNSQTFDYDIALLQLSIA CTWKFQTSLSTLGIALKFYNYSITKKSMKGCEHGWWEINEHMYCGSYMDHQTIFRVPSPLVHIQLQCSSRLSDKPLLA SSILYRICEPTRTLMSFVSTNNLMLVTFKSPHIRRLSGIRAYFEVIPEQKCENTVLVKDITGFEGKISSPYYPSYYPPKCK EQNCTQSIPCNNRTFKCGNDICFRKQNAKCDGTVDCPDGSDEEGCTCSRSSSALHRIIGGTDTLEGGWPWQVSLHF NPETLKQLIQPICIPPTGQRVRSGEKCWVTGWGRRHEADNKGSLVLQQAEVELIDQTLCVSTYGIITSRMLCAGIMSG **-TVFLFILAVIAWTLLWLYISKTESKDAFYFAGMFRITNIEFLPEYRQKESREFLSVSRTVQQVINLVYTTSAFSKFYEQS** KRDACKGDSGGPLSCRRKSDGKWILTGIVSWGHGCGRPNFPGVYTRVSNFVPWIHKYVPSL

>SGPr422 SEQID 109

DDIRQICLPEASASFQPNLTVHITGFGALYYGGESQNDLREARVKIISDDVCKQPQVYGNDIKPGMFCAGYMEGIYDAC MTLNKIKDLFAGKGQWDLAPEAEMLKPWMIAVLIVLSLTVVAVTIGLLVHFLVFDQKKEYYHGSFKILDPQINNNFGQSN DNIHQCGATLISNTWLVTAAHCFQKYKNPHQWTVSFGTKINPPLMKRNVRRFIIHEKYRSAAREYDIAVVQVSSRVTFS TYQLKDLRETTENLVYSLKMYLSFVCHSPEEDGVKVDVIMVFQFPSTEQRAVREKKIQSILNQKIRNLRALPINASSVQV NVAMVKNGNVGPGSGAGEAPGLGAGPAWSPMSSSTGELTVQASCGKRVVPLNVNRIASGVIAPKAAWPWQASLQY RGDSGGPLVTRDLKDTWYLIGIVSWGDNCGQKDKPGVYTQVTYYRNWIASKTGI

>SGPr538\_SEQID\_110

YDVALLRLQTALNFSDTVGAVCLPAKEQHFPKGSRCWVSGWGHTHPSHTYSSDMLQDTVVPLFSTQLCNSSCVYSG ASQPISGTLQDEEITLSCSEASAEEALLPALPKTVSFRINSEDFLLEAQVRDQPRWLLVCHEGWSPALGLQICWSLGHL NQASVALGFRHTCGGSVLAPRWVVTAAHCMHSFRLARLSSWRVHAGLVSHSAVRPHQGALVERIIPHPLYSAQNHD RLTHHKGVNLTDIKLNSSQEFAQLSPRLGGFLEEAWQPRNNCTSGQVVSLRCSECGARPLASRIVGGQSVAPGRWP MSLMLDDQPPMEAQYAEEGPGPGIFRAEPGDQQHPISQAVCWRSMRRGCAVLGALGLLAGAGVGSWLLVLYLCPA 4LTPRMLCAGYLDGRADACQGDSGGPLVCPDGDTWRLVGVVSWGRACAEPNHPGVYAKVAEFLDWIHDTAQDSLI

>SGPr527 1 SEQID 111

/LGRAGASSLPQGHQVSRLVISIRLPQHLGLRPPLALLELSSRVEPSPSALPICLHPAGIPPGASCWVLGWKEPQDRVP MARHLLLPLVMLVISPIPGAFQDSALSPTQEEPEDLDCGRPEPSARIVGGSNAQPGTWPWQVSLHHGGGHICGGSLIA **VAAAVSILTQRICDCLYQGILPPGTLCVLYAEGQENRCEMTSAPPLLCQMTEGSWILVGMAVQGSRELFAAIGPEEAWI** WPVCLPRASHRFVHGTACWATGWGDVQEADPLPLPWVLQEVELRLLGEATCQCLYSQPGPFNLTLQILPGMLCAGY QSDPQEPREENCTIALPECGKAPRPGAWPWEAQVMVPGSRPCHGALVSESWVLAPASCFLDPNSSDSPPRDLDAW PEGRRDTCQGDSGGPLVCEEGGRWFQAGITSFGFGCGRRNRPGVFTAVATYEAWIREQVMGSEPGPAFPTQPQKT PGALLEAELLGGWWCHCLYGRQGAAVPLPGDPPHALCPAYQEKEEVGSCWTHGPWISHVTRGAYLEDQLAWDWG PDGEETETQTCPPHTEHGACGLRLEAAPVGVLWPWLAEVHVAGDRVCTGILLAPGWVLAATHCVLRPGSTTVPYIEV 28WVLSAAHCFMTNGTLEPAAEWSVLLGVHSQDGPLDGAHTRAVAAIVVPANYSQVELGADLALLRLASPASLGPAV RVLLPSRPRAERVARLVQHENASWDNASDLALLQLRTPVNLSAASRPVCLPHPEHYFLPGSRCRLARWGRGEPALG SQTVGEANFLPPSGSPHWPTGGSNLCPPELAKASGSPHAVYFLLLLTLLIQS

SGPr542\_SEQID\_112

4GTRCRVAGWGFVSDFEELPPGLMEAKVRVLDPDVCNSSWKGHLTLTMLCTRSGDSHRRGFCSADSGGPLVCRNR **CFSHRDLRTGLVVLGAHVLSTAEPTQQVFGIDALTTHPDYHPMTHANDICLLQLNGSAVLGPAVGLLRLPGRRARPPT** AMGLGLRGWGRPLLTVATALMLPVKPPAGSWGAQIIGGHEVTPHSRPYMASVRFGGQHHCGGFLLRARWVVSAAH AHGLVSFSGLWCGDPKTPDVYTQVSAFVAWIWDVVRRSSPQPGPLPGTTRPPGEAA

>SGPr551 SEQID 113

MLSPEVVQALLVEELLSTVNSSAAVPYRAEYEVDPEGLVILEASVKDIAALNSTLGCYRYSYVGQGQVLRLKGPDHLAS JPFVLSVQPVVFQACEVNLTLDNRLDSQGVLSTPYFPSYYSPQTHCSWHLTVPSLDYGLALWFDAYALRRQKYDLPC TOGOWTIONRRLCGLRILQPYAERIPVVATAGITINFTSQISLTGPGVRVHYGLYNQSDPCPGEFLCSVNGLCVPACDG WPVAEAPQVAGGGGGGGGGGEEAEPEGMFKACEDSKRKARGYLRLVPLFVLLALLVLASAGVLLWYFLGYKAEVMVS SCLWHLQGPKDLMLKLRLEWTLAECRDRLAMYDVAGPLEKRLITSVYGCSRQEPVVEVLASGAIMAVVWKKGLHSYY WITGWGALREGGPISNALQKVDVQLIPQDLCSEAYRYQVTPRMLCAGYRKGKKDACQGDSGGPLVCKALSGRWFLA QVYSGSLRVLNRHFSQDLTRRESSAFRSETAKAQKMLKELITSTRLGTYYNSSSVYSFGEGPLTCFFWFILQIPEHRRL **JGRPDCRDGSDEEHCDCGLQGPSSRIVGGAVSSEGEWPWQASLQVRGRHICGGALIADRWVITAAHCFQEDSMAS** /KDCPNGLDERNCVCRATFQCKEDSTCISLPKVCDGQPDCLNGSDEEQCQEGVPCGTFTFQCEDRSCVKKPNPQC VLWTVFLGKVWQNSRWPGEVSFKVSRLLLHPYHEEDSHDYDVALLQLDHPVVRSAAVRPVCLPARSHFFEPGLHC 3LVSWGLGCGRPNYFGVYTRITGVISWIQQVVT >SGPr451 SEQID 114

TAPLKDVLQGSRIIGGTEAQAGAWPWVVSLQIKYGRVLVHVCGGTLVRERWVLTAAHCTKDTSDPLMWTAVIGTNNIH AEVHYISREMCNSERSYGGIIPNTSFCAGDEDGAFDTCRGDSGGPLMCYLPEYKRFFVMGITSYGHGCGRRGFPGVY DLPPSCSPASKMRLGLLSVALLFVGSSHLYSDHYSPSGRHRLGPSPEPAASSQQAEAVRKRLRRREGGAHAKDCG GRYPHTKKIKIKAIIIHPNFILESYVNDIALFHLKKAVRYNDYIQPICLPFDVFQILDGNTKCFISGWGRTKEEGNATNILQD **GPSFYQKWLTEHFFHASTQGILTINILRGQILIALCFVILLATT** 

>SGPr452 1 SEQID 115

GHNAPPGKWPWQVSLRVYSYHWASWAHICGGSLIHPQWVLTAAHCIFWKDTDPSIYRIHAGDVYLYGGRGLLNVSRI SPPQPRTPDCRLQASLEALATLAPQPSDWLCFADLGWFEADGAAHSMGLGSSLKWAWAKPSGMPVPENDLVGIVG VHPNYVTAGLGADVALLQLPGSPLSPESLPPPYRLQQASVQVLENAVCEQPYRNASGHTGDRQLILDDMLCAGSEG R D S C Y G D S G G P L V C R L R G S W R L V G V S W G Y G C T L R D F P G V Y T H V Q I Y V L W I L Q Q V G E L P

>SGPr504\_SEQID\_116 IIGGHEVTPHSRPYMASVRFGGQHHCGGFLLRARWVVSAAQCFSH >SGPr469\_SEQID\_117 GDSGGPLVCELNGTWVQVGIVSWGIGCGRKGYPGVYTEVSFYKKWI

>SGPr400\_SEQID\_118

HVCSDTLISEEWVLTVAICFPLSPHPDFQANTSSAIAVVELPSPVSVSPVVLLICLPSSEVYLKKNTTSCWVTGWGYTGI \_KSGPFRIWQGVKTKGEEGDRDTGTAGYAFTLLLLLGISGEPPEWVCGRPTVSSGIASGLGASVGQWPWQVSIRQGI <u>MAGEQVTANVSRYPGQKTMSFPEKTFLLSYRASLLAVVTHRSNNSRGRAFESQVLPDLTAGDAADPPIPPLGPGAAI</u> FQYIKRSYTLKELKVPLIDLQTCGDHYQNEILLHGVELIISEAMICSKLPVGQMDQCTVRIHPSGTFHRPCLPQ